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10/501834

## SEQUENCE LISTING

<110> Harris, Peter C., Ward, Christopher J., Rossetti, Sandro, and Torres, Vicente E.

<120> Polycystic Kidney Disease Nucleic Acids  
and Proteins

<130> 07039/386US1

<140> 10/501,834

<141> 2004-07-19

<150> PCT/US03/02038

<151> 2003-01-23

<150> 60/351,110

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agtggcagct tggggctttc ccaagagaag aaagcctcct gcggggccac tgaggcatc 12180
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<210> 2
<211> 4074
<212> PRT
<213> Homo sapien

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<400> 2
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Ala Val Arg His Leu Ser Leu His Ile Glu Pro Glu Glu Gly Ser Leu
20          25          30
Ala Gly Gly Thr Trp Ile Thr Val Ile Phe Asp Gly Leu Glu Leu Gly
35          40          45
Val Leu Tyr Pro Asn Asn Gly Ser Gln Leu Glu Ile His Leu Val Asn
50          55          60
Val Asn Met Val Val Pro Ala Leu Arg Ser Val Pro Cys Asp Val Phe
65          70          75          80
Pro Val Phe Leu Asp Leu Pro Val Val Thr Cys Arg Thr Arg Ser Val
85          90          95
Leu Ser Glu Ala His Glu Gly Leu Tyr Phe Leu Glu Ala Tyr Phe Gly
100         105         110
Gly Gln Leu Val Ser Ser Pro Asn Pro Gly Pro Arg Asp Ser Cys Thr
115         120         125
Phe Lys Phe Ser Lys Ala Gln Thr Pro Ile Val His Gln Val Tyr Pro

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130	135	140
Pro Ser Gly Val	Pro Gly Lys Leu Ile His	Val Tyr Gly Trp Ile Ile
145	150	155
Thr Gly Arg Leu	Glu Thr Phe Asp Phe Asp	Ala Glu Tyr Ile Asp Ser
165	170	175
Pro Val Ile Leu	Glu Ala Gln Gly Asp Lys	Trp Val Thr Pro Cys Ser
180	185	190
Leu Ile Asn Arg	Gln Met Gly Ser Cys Tyr	Pro Ile Gln Glu Asp His
195	200	205
Gly Leu Gly Thr	Leu Gln Cys His Val Glu	Gly Asp Tyr Ile Gly Ser
210	215	220
Gln Asn Val Ser	Phe Ser Val Phe Asn Lys	Gly Lys Ser Met Val His
225	230	235
Lys Lys Ala Trp	Leu Ile Ser Ala Lys Gln	Asp Leu Phe Leu Tyr Gln
245	250	255
Thr His Ser Glu	Ile Leu Ser Val Phe Pro	Glu Thr Gly Ser Leu Gly
260	265	270
Gly Arg Thr Asn	Ile Thr Ile Thr Gly Asp	Phe Phe Asp Asn Ser Ala
275	280	285
Gln Val Thr Ile	Ala Gly Ile Pro Cys Asp	Ile Arg His Val Ser Pro
290	295	300
Arg Lys Ile Glu	Cys Thr Thr Arg Ala Pro	Gly Lys Asp Val Arg Leu
305	310	315
Thr Thr Pro Gln	Pro Gly Asn Arg Gly Leu	Leu Phe Glu Val Gly Asp
325	330	335
Ala Val Glu Gly	Leu Glu Leu Thr Glu Ala	Thr Pro Gly Tyr Arg Trp
340	345	350
Gln Ile Val Pro	Asn Ala Ser Ser Pro Phe	Gly Phe Trp Ser Gln Glu
355	360	365
Gly Gln Pro Phe	Arg Ala Arg Leu Ser Gly	Phe Phe Val Ala Pro Glu
370	375	380
Thr Asn Asn Tyr	Thr Phe Trp Ile Gln Ala	Asp Ser Gln Ala Ser Leu
385	390	395
His Phe Ser Trp	Ser Glu Glu Pro Arg Thr	Lys Val Lys Val Ala Ser
405	410	415
Ile Ser Val Gly	Thr Ala Asp Trp Phe Asp	Ser Trp Glu Gln Asn Arg
420	425	430
Asp Glu Gly Thr	Trp Gln Gln Lys Thr Pro	Lys Leu Glu Leu Leu Gly
435	440	445
Gly Ala Met Tyr	Tyr Leu Glu Ala Glu His	His Gly Ile Ala Pro Ser
450	455	460
Arg Gly Met Arg	Ile Gly Val Gln Ile His	Asn Thr Trp Leu Asn Pro
465	470	475
Asp Val Val Thr	Thr Tyr Leu Arg Glu Lys	His Gln Ile Arg Val Arg
485	490	495
Ala Gln Arg Leu	Pro Glu Val Gln Val Leu	Asn Val Ser Gly Arg Gly
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Asn Phe Phe Leu	Thr Trp Asp Asn Val Ser	Ser Ser Gln Pro Ile Pro Ala
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Asn Ala Thr Ala	His Leu Ile Gln Thr Thr	Ile Glu Leu Leu Ala
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Val Lys Cys Lys	Leu Glu Pro Leu Trp Ser	Asn Ile Leu Leu Arg Leu

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Gly	Phe	Glu	Arg	Gly	Pro	Glu	Val	Ser	Asn	Ser	Asp	Gly	Asp	Leu	Thr
				565						570				575	
Ser	Gly	Thr	Glu	Pro	Phe	Cys	Gly	Arg	Phe	Ser	Leu	Arg	Gln	Pro	Arg
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His	Leu	Val	Leu	Thr	Pro	Pro	Ala	Ala	Gln	Lys	Gly	Tyr	Arg	Leu	Asp
		595					600					605			
Gln	Tyr	Thr	His	Leu	Cys	Leu	Ala	Tyr	Lys	Gly	His	Met	Asn	Lys	Ile
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Leu	Lys	Met	Ile	Val	Ser	Phe	Thr	Ile	Gly	Phe	Gln	Asn	Met	Val	Lys
625					630					635				640	
Asn	Thr	Thr	Cys	Asp	Trp	Ser	Leu	Thr	Arg	Thr	Ser	Pro	Glu	Ser	Trp
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Gln	Phe	Asp	Cys	Thr	Asp	Leu	Trp	Glu	Thr	Cys	Val	Arg	Cys	Phe	Gly
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Asp	Leu	Gln	Pro	Pro	Pro	Ala	Asn	Ser	Pro	Val	Leu	Val	His	Gln	Ile
	675						680					685			
Asn	Leu	Leu	Pro	Leu	Ala	Gln	Glu	Thr	Gly	Leu	Phe	Tyr	Val	Asp	Glu
	690					695					700				
Ile	Ile	Ile	Ala	Asp	Thr	Asn	Val	Thr	Val	Ser	Gln	Ala	Asp	Ser	Gly
705					710					715				720	
Thr	Ala	Arg	Pro	Gly	Gly	Asn	Leu	Val	Glu	Ser	Val	Ser	Val	Val	Gly
			725						730					735	
Ser	Pro	Pro	Val	Tyr	Ser	Val	Thr	Ser	Trp	Leu	Ala	Gly	Cys	Gly	Thr
		740					745					750			
Glu	Leu	Pro	Leu	Ile	Thr	Ala	Arg	Ser	Val	Pro	Thr	Glu	Gly	Thr	Glu
	755					760						765			
Glu	Gly	Ser	Gly	Leu	Val	Leu	Val	Thr	Thr	Gln	Arg	Arg	Gln	Arg	Thr
	770					775				780					
Ser	Pro	Pro	Leu	Gly	Gly	His	Phe	Arg	Ile	Gln	Leu	Pro	Asn	Thr	Val
785					790					795				800	
Ile	Ser	Asp	Val	Pro	Val	Gln	Ile	Ser	Ala	His	His	Leu	His	Gln	Leu
			805						810					815	
Leu	Gln	Asn	Asn	Ala	Asp	Asp	Phe	Thr	Ser	Arg	Tyr	Leu	Asn	Ala	Ser
		820						825					830		
Asp	Phe	Thr	Val	Lys	Glu	Asp	Leu	Tyr	Thr	Cys	Tyr	Glu	His	Val	Trp
	835						840					845			
Thr	Leu	Ser	Trp	Ser	Thr	Gln	Ile	Gly	Asp	Leu	Pro	Asn	Phe	Ile	Arg
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Val	Ser	Asp	Glu	Asn	Leu	Thr	Gly	Val	Asn	Pro	Ala	Ala	Ala	Thr	Arg
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Val	Val	Tyr	Asp	Gly	Gly	Val	Phe	Leu	Gly	Pro	Ile	Phe	Gly	Asp	Met
			885						890					895	
Leu	Ala	Thr	Ala	Asn	Gln	His	Thr	Gln	Val	Val	Val	Arg	Val	Asn	Asp
		900						905					910		
Val	Pro	Ala	His	Cys	Pro	Gly	Ser	Cys	Ser	Phe	Gln	Tyr	Leu	Gln	Gly
	915						920					925			
Ser	Thr	Pro	Cys	Val	His	Ser	Val	Trp	Tyr	Ser	Ile	Asp	Gly	Asp	Ile
	930					935					940				
Asn	Leu	Met	Ile	Tyr	Ile	Thr	Gly	Thr	Gly	Phe	Ser	Gly	Asp	Ser	Gln
945					950					955				960	
Phe	Leu	Gln	Val	Thr	Val	Asn	Lys	Thr	Ser	Cys	Lys	Val	Ile	Phe	Ser

				965					970					975
Asn	Gln	Thr	Asn	Val	Val	Cys	Gln	Thr	Asp	Leu	Leu	Pro	Val	Gly Met
			980					985					990	
His	Arg	Ile	Leu	Met	Leu	Val	Arg	Pro	Ser	Gly	Leu	Ala	Ile	Ser Ala
		995					1000					1005		
Thr	Gly	Glu	Asp	Leu	Phe	Leu	Asn	Val	Lys	Pro	Arg	Leu	Asp	Met Val
	1010					1015					1020			
Glu	Pro	Ser	Arg	Ala	Ala	Asp	Ile	Gly	Gly	Leu	Trp	Ala	Thr	Ile Arg
1025				1030						1035				1040
Gly	Ser	Ser	Leu	Glu	Gly	Val	Ser	Leu	Ile	Leu	Phe	Gly	Ser	Tyr Ser
			1045						1050					1055
Cys	Ala	Ile	Asn	Val	Ala	Thr	Ser	Asn	Ser	Ser	Arg	Ile	Gln	Cys Lys
			1060					1065					1070	
Val	Pro	Pro	Arg	Gly	Lys	Asp	Gly	Arg	Ile	Val	Asn	Val	Thr	Val Ile
	1075						1080					1085		
Arg	Gly	Asp	Tyr	Ser	Ala	Val	Leu	Pro	Arg	Ala	Phe	Thr	Tyr	Val Ser
	1090					1095					1100			
Ser	Leu	Asn	Pro	Val	Ile	Val	Thr	Leu	Ser	Arg	Asn	Ile	Ser	Asn Ile
1105				1110						1115				1120
Ala	Gly	Gly	Glu	Thr	Leu	Val	Ile	Gly	Val	Ala	Arg	Leu	Met	Asn Tyr
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Thr	Asp	Leu	Asp	Val	Glu	Val	His	Val	Gln	Asp	Ala	Leu	Ala	Pro Val
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His	Thr	Gln	Ser	Ala	Trp	Gly	Leu	Glu	Val	Ala	Leu	Pro	Pro	Leu Pro
	1155					1160						1165		
Ala	Gly	Leu	His	Arg	Ile	Ser	Val	Ser	Ile	Asn	Gly	Val	Ser	Ile His
	1170				1175						1180			
Ser	Gln	Gly	Val	Asp	Leu	His	Ile	Gln	Tyr	Leu	Thr	Glu	Val	Phe Ser
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Ile	Glu	Pro	Cys	Cys	Gly	Ser	Leu	Leu	Gly	Gly	Thr	Ile	Leu	Ser Ile
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Ser	Gly	Ile	Gly	Phe	Ser	Arg	Asp	Pro	Ala	Leu	Val	Trp	Val	Leu Val
	1220							1225					1230	
Gly	Asn	Arg	Ser	Cys	Asp	Ile	Val	Asn	Leu	Thr	Glu	Ala	Ser	Ile Trp
	1235						1240					1245		
Cys	Glu	Thr	Leu	Pro	Ala	Pro	Gln	Ile	Pro	Asp	Ala	Gly	Ala	Pro Thr
	1250					1255					1260			
Val	Pro	Ala	Ala	Val	Glu	Val	Trp	Ala	Gly	Asn	Arg	Phe	Phe	Ala Arg
1265				1270						1275				1280
Gly	Pro	Ser	Pro	Ser	Leu	Val	Gly	Lys	Gly	Phe	Thr	Phe	Met	Tyr Glu
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Ala	Ala	Ala	Thr	Pro	Val	Val	Thr	Ala	Met	Gln	Gly	Glu	Ile	Thr Asn
	1300							1305					1310	
Ser	Ser	Leu	Ser	Leu	His	Val	Gly	Gly	Ser	Asn	Leu	Ser	Asn	Ser Val
	1315						1320					1325		
Ile	Leu	Leu	Gly	Asn	Leu	Asn	Cys	Asp	Val	Glu	Thr	Gln	Ser	Phe Gln
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Ala	Gly	Ile	Tyr	Pro	Leu	Gln	Val	Arg	Gln	Lys	Gln	Met	Gly	Phe Ala
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Asn	Met	Ser	Val	Val	Leu	Gln	Gln	Phe	Ala	Val	Met	Pro	Arg	Ile Met



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Ala Ile Phe Pro Ser Gln Gly Ser Ala Cys Gly Gly Thr Ile Leu Thr		
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Val Arg Gly Leu Leu Leu Asn Ser Arg Arg Arg Ser Val Arg Val Asp		
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Leu Ser Gly Pro Phe Thr Cys Val Ile Leu Ser Leu Gly Asp His Thr		
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Ile Leu Cys Gln Val Ser Leu Glu Gly Asp Pro Leu Pro Gly Ala Ser		1440
1445	1450	1455
Phe Ser Leu Asn Val Thr Val Leu Val Asn Gly Leu Thr Ser Glu Cys		
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Gln Gly Asn Cys Thr Leu Phe Ile Arg Glu Glu Ala Ser Pro Val Met		
1475	1480	1485
Asp Ala Leu Ser Thr Asn Thr Ser Gly Ser Leu Thr Thr Val Leu Ile		
1490	1495	1500
Arg Gly Gln Arg Leu Ala Thr Thr Ala Asp Glu Pro Met Val Phe Val		
1505	1510	1515
Asp Asp Gln Leu Pro Cys Asn Val Thr Phe Phe Asn Ala Ser His Val		1520
1525	1530	1535
Val Cys Gln Thr Arg Asp Leu Ala Pro Gly Pro His Tyr Leu Ser Val		
1540	1545	1550
Phe Tyr Thr Arg Asn Gly Tyr Ala Cys Ser Gly Asn Val Ser Arg His		
1555	1560	1565
Phe Tyr Ile Met Pro Gln Val Phe His Tyr Phe Pro Lys Asn Phe Ser		
1570	1575	1580
Leu His Gly Gly Ser Leu Leu Thr Ile Glu Gly Thr Gly Leu Arg Gly		
1585	1590	1595
Gln Asn Thr Thr Ser Val Tyr Ile Asp Gln Gln Thr Cys Leu Thr Val		1600
1605	1610	1615
Asn Ile Gly Ala Glu Leu Ile Arg Cys Ile Val Pro Thr Gly Asn Gly		
1620	1625	1630
Ser Val Ala Leu Glu Ile Glu Val Asp Gly Leu Trp Tyr His Ile Gly		
1635	1640	1645
Val Ile Gly Tyr Asn Lys Ala Phe Thr Pro Glu Leu Ile Ser Ile Ser		
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Gln Ser Asp Asp Ile Leu Thr Phe Ala Val Ala Gln Ile Ser Gly Ala		
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Ala Asn Ile Asp Ile Phe Ile Gly Met Ser Pro Cys Val Gly Val Ser		1680
1685	1690	1695
Gly Asn His Thr Val Leu Gln Cys Val Val Pro Ser Leu Pro Ala Gly		
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Glu Tyr His Val Arg Gly Tyr Asp Cys Ile Arg Gly Trp Ala Ser Ser		
1715	1720	1725
Ala Leu Val Phe Thr Ser Arg Val Ile Ile Thr Ala Val Thr Glu Asn		
1730	1735	1740
Phe Gly Cys Leu Gly Gly Arg Leu Val His Val Phe Gly Ala Gly Phe		
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Ser Pro Gly Asn Val Ser Ala Ala Val Cys Gly Ala Pro Cys Arg Val		1760
1765	1770	1775
Leu Ala Asn Ala Thr Val Ser Ala Phe Ser Cys Leu Val Leu Pro Leu		
1780	1785	1790
Asp Val Ser Leu Ala Phe Leu Cys Gly Leu Lys Arg Glu Glu Asp Ser		

1795	1800	1805
Cys Glu Ala Ala Arg His Thr Tyr Val Gln Cys Asp Leu Thr Val Ala		
1810	1815	1820
Met Ala Thr Glu Gln Leu Leu Glu Ser Trp Pro Tyr Leu Tyr Ile Cys		
1825	1830	1835
Glu Glu Ser Ser Gln Cys Leu Phe Val Pro Asp His Trp Ala Glu Ser		1840
1845	1850	1855
Met Phe Pro Ser Phe Ser Gly Leu Phe Ile Ser Pro Lys Leu Glu Arg		
1860	1865	1870
Asp Glu Val Leu Ile Tyr Asn Ser Ser Cys Asn Ile Thr Met Glu Thr		
1875	1880	1885
Glu Ala Glu Met Glu Cys Glu Thr Pro Asn Gln Pro Ile Thr Val Lys		
1890	1895	1900
Ile Thr Glu Ile Arg Lys Arg Trp Gly Gln Asn Thr Gln Gly Asn Phe		
1905	1910	1915
Ser Leu Gln Phe Cys Arg Arg Trp Ser Arg Thr His Ser Trp Phe Pro		1920
1925	1930	1935
Glu Arg Leu Pro Gln Asp Gly Asp Asn Val Thr Val Glu Asn Gly Gln		
1940	1945	1950
Leu Leu Leu Asp Thr Asn Thr Ser Ile Leu Asn Leu Leu His Ile		
1955	1960	1965
Lys Gly Gly Lys Leu Ile Phe Met Ala Pro Gly Pro Ile Glu Leu Arg		
1970	1975	1980
Ala His Ala Ile Leu Val Ser Asp Gly Gly Glu Leu Arg Ile Gly Ser		
1985	1990	1995
Glu Asp Lys Pro Phe Gln Gly Arg Ala Gln Ile Thr Leu Tyr Gly Ser		2000
2005	2010	2015
Ser Tyr Ser Thr Pro Phe Phe Pro Tyr Gly Val Lys Phe Leu Ala Val		
2020	2025	2030
Arg Asn Gly Thr Leu Ser Leu His Gly Ser Leu Pro Glu Val Ile Val		
2035	2040	2045
Thr Cys Leu Arg Ala Thr Ala His Ala Leu Asp Thr Val Leu Ala Leu		
2050	2055	2060
Glu Asp Ala Val Asp Trp Asn Pro Gly Asp Glu Val Val Ile Ile Ser		
2065	2070	2075
Gly Thr Gly Val Lys Gly Ala Lys Pro Met Glu Glu Ile Val Thr Val		2080
2085	2090	2095
Glu Thr Val Gln Asp Thr Asp Leu Tyr Leu Lys Ser Pro Leu Arg Tyr		
2100	2105	2110
Ser His Asn Phe Thr Glu Asn Trp Val Ala Gly Glu His His Ile Leu		
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Lys Ala Thr Val Ala Leu Leu Ser Arg Ser Ile Thr Ile Gln Gly Asn		
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Leu Thr Asn Glu Arg Glu Lys Leu Leu Val Ser Cys Gln Glu Ala Asn		
2145	2150	2155
Ala Pro Glu Gly Asn Leu Gln His Cys Leu Tyr Ser Met Ser Glu Lys		2160
2165	2170	2175
Met Leu Gly Ser Arg Asp Met Gly Ala Arg Val Ile Val Gln Ser Phe		
2180	2185	2190
Pro Glu Glu Pro Ser Gln Val Gln Leu Lys Gly Val Gln Phe Gln Val		
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Leu Gly Gln Ala Phe His Lys His Leu Ser Ser Leu Thr Leu Val Gly		

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Ala Met Arg Glu Ser Phe	Ile Gln Gly Cys Thr Val Arg Asn Ser Phe	
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Ser Arg Gly Leu Ser Met Cys Gly Thr Leu Gly Leu Lys Val Asp Ser		2240
2245	2250	2255
Asn Val Phe Tyr Asn Ile Leu Gly His Ala Leu Leu Val Gly Thr Cys		
2260	2265	2270
Thr Glu Met Arg Tyr Ile Ser Trp Glu Ala Ile His Gly Arg Lys Asp		
2275	2280	2285
Asp Trp Ser Gly His Gly Asn Ile Ile Arg Asn Asn Val Ile Ile Gln		
2290	2295	2300
Val Ser Gly Ala Glu Gly Leu Ser Asn Pro Glu Met Leu Thr Pro Ser		
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Gly Ile Tyr Ile Cys Ser Pro Thr Asn Val Ile Glu Gly Asn Arg Val		2320
2325	2330	2335
Cys Gly Ala Gly Tyr Gly Tyr Phe Phe His Leu Met Thr Asn Gln Thr		
2340	2345	2350
Ser Gln Ala Pro Leu Leu Ser Phe Thr Gln Asn Ile Ala His Ser Cys		
2355	2360	2365
Thr Arg Tyr Gly Leu Phe Val Tyr Pro Lys Phe Gln Pro Pro Trp Asp		
2370	2375	2380
Asn Val Thr Gly Thr Thr Leu Phe Gln Ser Phe Thr Val Trp Glu Ser		
2385	2390	2395
Ala Gly Gly Ala Gln Ile Phe Arg Ser Ser Asn Leu Arg Leu Lys Asn		
2405	2410	2415
Phe Lys Val Tyr Ser Cys Arg Asp Phe Gly Ile Asp Val Leu Glu Ser		
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Asp Ala Asn Thr Ser Val Thr Asp Ser Leu Leu Leu Gly His Phe Ala		
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His Lys Gly Ser Leu Cys Met Ser Ser Gly Ile Lys Thr Pro Lys Arg		
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Trp Glu Leu Met Val Ser Asn Thr Thr Phe Val Asn Phe Asp Leu Ile		
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Asn Cys Val Ala Ile Arg Thr Cys Ser Asp Cys Ser Gln Gly Gln Gly		2480
2485	2490	2495
Gly Phe Thr Val Lys Thr Ser Gln Leu Lys Phe Thr Asn Ser Ser Asn		
2500	2505	2510
Leu Val Ala Phe Pro Phe Pro His Ala Ala Ile Leu Glu Asp Leu Asp		
2515	2520	2525
Gly Ser Leu Ser Gly Lys Asn Arg Ser His Ile Leu Ala Ser Met Glu		
2530	2535	2540
Thr Leu Ser Ala Ser Cys Leu Val Asn Ser Ser Phe Gly Arg Val Val		
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His Gly Ser Ala Cys Gly Gly Gly Val Leu Phe His Arg Met Ser Ile		2560
2565	2570	2575
Gly Leu Ala Asn Thr Pro Glu Val Ser Tyr Asp Leu Thr Met Thr Asp		
2580	2585	2590
Ser Arg Asn Lys Thr Thr Thr Val Asn Tyr Val Arg Asp Thr Leu Ser		
2595	2600	2605
Asn Pro Arg Gly Trp Met Ala Leu Leu Leu Asp Gln Glu Thr Tyr Ser		
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Leu Gln Ser Glu Asn Leu Trp Ile Asn Arg Ser Leu Gln Tyr Ser Ala		

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Thr Phe Asp Asn Phe Ala Pro Gly Asn Tyr Leu Leu Leu Val His Thr						
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Asp Leu Pro Pro Tyr Pro Asp Ile Leu Leu Arg Cys Gly Ser Arg Val						
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Gly Leu Ser Phe Pro Phe Leu Pro Ser Pro Gly Gln Asn Gln Gly Cys						
	2675		2680		2685	
Asp Trp Phe Phe Asn Ser Gln Leu Arg Gln Leu Thr Tyr Leu Val Ser						
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Gly Glu Gly Gln Val Gln Val Ile Leu Arg Val Lys Glu Gly Met Pro						
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Pro Thr Ile Ser Ala Ser Thr Ser Ala Pro Glu Ser Ala Leu Lys Trp						
	2725		2730		2735	
Ser Leu Pro Glu Thr Trp Gln Gly Val Glu Glu Gly Trp Gly Gly Tyr						
	2740		2745		2750	
Asn Asn Thr Ile Pro Gly Pro Gly Asp Asp Val Leu Ile Leu Pro Asn						
	2755		2760		2765	
Arg Thr Val Leu Val Asp Thr Asp Leu Pro Phe Phe Lys Gly Leu Tyr						
	2770		2775		2780	
Val Met Gly Thr Leu Asp Phe Pro Val Asp Arg Ser Asn Val Leu Ser						
2785		2790		2795		2800
Val Ala Cys Met Val Ile Ala Gly Gly Glu Leu Lys Val Gly Thr Leu						
	2805		2810		2815	
Glu Asn Pro Leu Glu Lys Glu Gln Lys Leu Leu Ile Leu Leu Arg Ala						
	2820		2825		2830	
Ser Glu Gly Val Phe Cys Asp Arg Met Asn Gly Ile His Ile Asp Pro						
	2835		2840		2845	
Gly Thr Ile Gly Val Tyr Gly Lys Val His Leu Tyr Ser Ala Tyr Pro						
	2850		2855		2860	
Lys Asn Ser Trp Thr His Leu Gly Ala Asp Ile Ala Ser Gly Asn Glu						
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Arg Ile Ile Val Glu Asp Ala Val Asp Trp Arg Pro His Asp Lys Ile						
	2885		2890		2895	
Val Leu Ser Ser Ser Tyr Glu Pro His Glu Ala Glu Val Leu Thr						
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Val Lys Glu Val Lys Gly His His Val Arg Ile Tyr Glu Arg Leu Lys						
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His Arg His Ile Gly Ser Val His Val Thr Glu Asp Gly Arg His Ile						
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Arg Leu Ala Ala Glu Val Gly Leu Leu Thr Arg Asn Ile Gln Ile Gln						
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Pro Asp Val Ser Cys Arg Gly Arg Leu Phe Val Gly Ser Phe Arg Lys						
	2965		2970		2975	
Ser Ser Arg Glu Glu Phe Ser Gly Val Leu Gln Leu Leu Asn Val Glu						
	2980		2985		2990	
Ile Gln Asn Phe Gly Ser Pro Leu Tyr Ser Ser Val Glu Phe Ser Asn						
	2995		3000		3005	
Val Ser Ala Gly Ser Trp Ile Ile Ser Ser Thr Leu His Gln Ser Cys						
	3010		3015		3020	
Gly Gly Gly Ile His Ala Ala Ala Ser His Gly Val Leu Leu Asn Asp						
3025		3030		3035		3040
Asn Ile Val Phe Gly Thr Ala Gly His Gly Ile Asp Leu Glu Gly Gln						

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Ala	Tyr	Thr	Val	Thr	Asn	Asn	Leu	Val	Val	Leu	Met	Thr	Gln	Pro	Ala
3060				3065				3070							
Trp	Ser	Thr	Ile	Trp	Val	Ala	Gly	Ile	Lys	Val	Asn	Gln	Val	Lys	Asp
3075				3080				3085							
Ile	Asn	Leu	His	Gly	Asn	Val	Val	Ala	Gly	Ser	Glu	Arg	Leu	Gly	Phe
3090				3095				3100							
His	Ile	Arg	Gly	His	Lys	Cys	Ser	Ser	Cys	Glu	Leu	Leu	Trp	Ser	Asp
3105					3110					3115					3120
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65          70          75          80
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Gly	Ser	Trp	His	Gln	Lys	Thr	Thr	Lys	Leu	Glu	Leu	Gln	Gly	Gly	Ala	435	440	445
Lys	Tyr	Tyr	Leu	Glu	Ala	Glu	Gln	His	Gly	Ile	Ala	Pro	Ser	Arg	Gly	450	455	460
Met	Arg	Ile	Gly	Val	Gln	Ile	His	Asn	Thr	Trp	Leu	Asn	Pro	Asp	Val	465	470	475
Val	Asn	Thr	Tyr	Leu	Leu	Glu	Lys	His	Gln	Ile	Arg	Ala	Arg	Ala	Gln	485	490	495
Arg	Leu	Pro	Glu	Ile	Gln	Val	Leu	His	Val	Ser	Gly	Lys	Gly	Asn	Phe	500	505	510
Phe	Leu	Thr	Trp	Gly	Asn	Val	Ser	Ser	Gln	Pro	Val	Pro	Ala	Asn	Ala	515	520	525
Thr	Ala	Gln	Gln	Ile	Gln	Thr	Thr	Ile	Glu	Glu	Leu	Leu	Val	Val	Lys	530	535	540

Cys	Asn	Leu	Ala	Pro	Phe	Ser	Ala	His	Val	Leu	Leu	Arg	Leu	Gly	Phe	545	550	555	560
Glu	Gln	Gly	Leu	Glu	Gly	Ser	Arg	Ser	Asp	Gly	Val	Arg	Thr	Ser	Ser	565	570	575	
Thr	Glu	Pro	Phe	Cys	Gly	Arg	Phe	Ser	Leu	Gly	Gln	Leu	Gly	His	Leu	580	585	590	
Ile	Leu	Ile	Pro	Glu	Ala	Ala	Asp	Lys	Gly	Tyr	Gln	Leu	Asp	Arg	Tyr	595	600	605	
Pro	Tyr	Leu	Cys	Leu	Ala	Tyr	Arg	Gly	His	Met	Asn	Lys	Thr	Leu	Asp	610	615	620	
Met	Thr	Val	Ser	Phe	Leu	Phe	Gly	Phe	Gln	Thr	Ile	Met	Lys	Asn	Ile	625	630	635	640
Thr	Cys	Asp	Trp	Ser	Leu	Thr	Asp	Pro	His	Pro	Glu	Ser	Trp	Gln	Phe	645	650	655	
Thr	Cys	Ile	Asn	Leu	Trp	Asp	Thr	Cys	Leu	Cys	His	Ser	Glu	Asp	Ile	660	665	670	
Gln	Ser	Ser	Leu	Ala	Asn	Thr	Pro	Leu	Leu	Ala	His	Arg	Ile	Asp	Ile	675	680	685	
Arg	Pro	Val	Val	Pro	Glu	Ala	Gly	Leu	Leu	Tyr	Val	Asp	Glu	Ile	Ile	690	695	700	
Leu	Ala	Asp	Thr	Asn	Val	Thr	Val	Ser	Gln	Ala	Asp	Ser	Gly	Arg	Ala	705	710	715	720
Cys	Pro	Gly	Gly	Asn	Val	Val	Glu	Ser	Val	Ser	Val	Val	Gly	Val	Pro	725	730	735	
Pro	Val	Tyr	Ser	Ile	Ser	Ser	Trp	Leu	Ala	Gly	Cys	Gly	Ser	Glu	Leu	740	745	750	
Pro	Leu	Ile	Thr	Ala	Cys	Ser	Val	Ser	Thr	Glu	Gly	Thr	Gly	Asp	Gly	755	760	765	
Ser	Glu	Leu	Ile	Glu	Val	Thr	Ala	Gln	Arg	Leu	Gln	Arg	Thr	Ser	Pro	770	775	780	
Pro	Leu	Gly	Gly	His	Phe	Phe	Leu	Tyr	Leu	Ser	Asp	Thr	Val	Ile	Pro	785	790	795	800
Asp	Val	Pro	Val	Arg	Met	Ser	Ala	Arg	Gln	Leu	His	Lys	Leu	Leu	Gln	805	810	815	
Asp	Ser	Ala	Asp	Glu	Ser	Thr	Ser	Gly	Tyr	Leu	Asn	Ala	Gly	Asp	Phe	820	825	830	
Thr	Val	Thr	Glu	Asp	Leu	Asn	Ser	Cys	Tyr	Glu	His	Val	Trp	Thr	Leu	835	840	845	
Ser	Trp	Thr	Thr	Gln	Thr	Gly	Asp	Leu	Pro	Asn	Phe	Ile	Arg	Val	Ser	850	855	860	
Asp	Gln	Asn	Leu	Thr	Gly	Val	Asn	Pro	Thr	Val	Thr	Ala	Arg	Val	Val	865	870	875	880
Tyr	Asp	Gly	Gly	Val	Phe	Leu	Gly	Pro	Ile	Phe	Gly	Asp	Met	Leu	Ala	885	890	895	
Thr	Ala	Asn	Gln	Gln	Thr	Gln	Val	Ala	Val	Gln	Val	Asn	Asp	Ile	Pro	900	905	910	
Ala	Tyr	Cys	Ser	Gly	Ser	Cys	Ser	Phe	Gln	Tyr	Gln	Gln	Glu	Ser	Thr	915	920	925	
Pro	Ser	Val	Asp	His	Val	Trp	Tyr	Ser	Leu	Gly	Ser	Asp	Val	Asn	Leu	930	935	940	
Leu	Val	His	Phe	Thr	Gly	Thr	Gly	Phe	Pro	Arg	Asp	Thr	Gln	Phe	Leu	945	950	955	960

Gln Val Thr Val Asn Lys Thr Ser Cys Glu Val Leu Phe Ser Asn Glu  
 965 970 975  
 Thr Asn Val Ala Cys Glu Leu Ala Leu Leu Pro Val Gly Val His Gln  
 980 985 990  
 Ile Phe Met Leu Val Ile Pro Ser Gly Leu Ala Val His Ala Ser Gly  
 995 1000 1005  
 Glu Asp Leu Leu Leu His Val Glu Pro Arg Leu Asp Ala Val Glu Pro  
 1010 1015 1020  
 Ser Thr Ala Ala Glu Ile Gly Gly Arg Trp Val Thr Leu Arg Gly Ser  
 1025 1030 1035 1040  
 Ser Leu Glu Gly Val Ser Leu Val Leu Phe Gly Thr Gln Ser Cys Val  
 1045 1050 1055  
 Ile Asp Ala Ile Arg Ser Asn Ser Gln Gln Ile Gln Cys Lys Val Pro  
 1060 1065 1070  
 Pro Arg Gly Lys Asp Gly Tyr Thr Val Asn Val Thr Val Ile Ser Gly  
 1075 1080 1085  
 Asp His Ser Thr Val Leu Ala Arg Ala Phe Thr Tyr Val Ser Ser Leu  
 1090 1095 1100  
 Asn Pro Val Ile Val Ser Leu Ser Arg Asn Arg Ser Ser Ile Ala Gly  
 1105 1110 1115 1120  
 Gly Glu Ile Leu Phe Leu Gly Met Ser Leu Leu Val Asn Tyr Thr Asp  
 1125 1130 1135  
 Leu Asp Val Gln Ile His Val Gln Asp Thr Ser Ala Gln Val Leu Ser  
 1140 1145 1150  
 Gln Thr Ala Trp Gly Leu Glu Val Val Leu Pro Pro Leu Val Pro Gly  
 1155 1160 1165  
 Ile His Val Ile Ser Ala Phe Ile Asn Gly Val Ser Ile Arg Ser Gln  
 1170 1175 1180  
 Gly Val Asp Leu Tyr Ile Gln Tyr Leu Thr Glu Val Phe Ser Val Glu  
 1185 1190 1195 1200  
 Pro Cys Ser Gly Ser Leu Leu Gly Val Phe Leu Leu Ser Leu Leu Arg  
 1205 1210 1215  
 Thr Gly Leu Gly Arg Asp Pro Ala Leu Ile Arg Val Leu Val Asp Asn  
 1220 1225 1230  
 His Pro Cys Asp Ile Val Asn Leu Thr Glu Val Asn Ile Trp Cys Glu  
 1235 1240 1245  
 Thr Pro Pro Ala Val Leu Pro Pro Arg Ala Asp Val Leu Thr Val Leu  
 1250 1255 1260  
 Ala Ser Val Glu Ile Trp Ala Gly Asn Thr Tyr Phe Phe His Gly Pro  
 1265 1270 1275 1280  
 Ser Leu Val Gly Lys Gly Phe Thr Phe Thr Tyr Glu Ala Ala Ala Thr  
 1285 1290 1295  
 Pro Val Val Thr Ala Met Trp Gly Glu Phe Lys Asn Asn Ser Val Arg  
 1300 1305 1310  
 Phe Tyr Val Glu Gly Ser Asn Ile Ser Asp Ser Val Ile Leu Leu Gly  
 1315 1320 1325  
 Ser Leu Lys Cys Glu Leu Glu Val Gln Phe Phe Gly Asp Ser Met Asn  
 1330 1335 1340  
 Leu Ser Gly Cys Phe Phe Pro Leu His Ser Leu Glu Ala Gly Val Tyr  
 1345 1350 1355 1360  
 Thr Leu Gln Val Arg His Lys Arg Met Gly Phe Ala Asn Met Ser Val  
 1365 1370 1375

Val Pro Gln Lys Phe Glu Leu Ser Pro Gln Ile Ile Ala Ile Phe Pro  
 1380 1385 1390  
 Thr His Gly Ser Lys Cys Gly Gly Thr Val Leu Thr Val Lys Gly Met  
 1395 1400 1405  
 Ala Phe Ser Ser Arg Lys Arg Ser Val His Val Asp Ile Ser Gly Pro  
 1410 1415 1420  
 Phe Ala Cys Met Ile Leu Ser Leu Glu Asp His Thr Val Leu Cys Gln  
 1425 1430 1435 1440  
 Thr Arg Phe Val Gly Asp Gln Phe Ser Glu Ala Ser Leu Ala Leu Asn  
 1445 1450 1455  
 Ile Thr Val Leu Val Asn Gly Leu Thr Ser Lys Cys Lys Gly Asn Cys  
 1460 1465 1470  
 Thr Leu Phe Ile Glu Glu Ala Ala Thr Pro Ile Val Asp Ala Leu Thr  
 1475 1480 1485  
 Ile Ser Ile Ser Gly Ser Leu Thr Met Val Leu Met Arg Gly Arg Arg  
 1490 1495 1500  
 Leu Ala Thr Thr Ala Asp Glu Pro Ile Ala Phe Val Asp Asp Gln Leu  
 1505 1510 1515 1520  
 Pro Cys His Thr Thr Phe Phe Asn Thr Ser His Val Ala Cys Gln Ile  
 1525 1530 1535  
 Arg Asp Leu Ala Pro Gly Phe His Tyr Leu Ser Ala Val His Thr Ser  
 1540 1545 1550  
 Ala Gly Tyr Ala Cys Leu Asn Ser Val Ser Arg Asn Phe Phe Ile Val  
 1555 1560 1565  
 Pro Gln Val Leu Asp Tyr Phe Pro Lys Asp Phe Ser Ile His Gly Gly  
 1570 1575 1580  
 Ser Leu Leu Thr Ile Lys Gly Thr Ala Leu Arg Gly Trp Lys Ala Thr  
 1585 1590 1595 1600  
 Val Val Tyr Val Gly Arg Gln Ala Cys Leu Thr Val Asn Phe Ser Ser  
 1605 1610 1615  
 Asp Phe Ile Gln Cys Ile Val Pro Ala Gly Asn Gly Ser Ala Ala Leu  
 1620 1625 1630  
 Glu Ile Asp Val Asn Gly Val Leu Tyr His Ile Gly Leu Val Asp Tyr  
 1635 1640 1645  
 Ser Ser Ile Phe Thr Pro Glu Leu Leu Ser Val Ser Arg Ser Gln Asp  
 1650 1655 1660  
 Ile Leu Thr Phe Thr Val Ala Arg Ile Ser Gly Ala Ala Asn Val Asp  
 1665 1670 1675 1680  
 Ile Phe Ile Gly Thr Ser Pro Cys Leu Gly Val Ala Gly Asn Arg Thr  
 1685 1690 1695  
 Val Leu Gln Cys Met Val Pro Leu Leu Pro Ala Gly Glu Tyr Leu Val  
 1700 1705 1710  
 Thr Gly Tyr Asp His Ser Arg Gly Trp Ala Ser Ser Thr Leu Ile Leu  
 1715 1720 1725  
 Val Leu Arg Ala Thr Val Thr Ser Val Thr Lys Asn Tyr Gly Cys Leu  
 1730 1735 1740  
 Gly Gly Arg Leu Leu His Val Leu Gly Ala Gly Phe Ser Pro Gly Asn  
 1745 1750 1755 1760  
 Ile Ser Ala Ala Val Cys Gly Ala Pro Cys Gln Val Leu Ala Asn Ala  
 1765 1770 1775  
 Thr Val Ser Ala Phe Ser Cys Leu Val Leu Pro Leu His Val Ser Leu  
 1780 1785 1790

Ala Phe Leu Cys Asp Leu Arg His Ala Glu Asp Ser Cys Lys Val Arg  
 1795 1800 1805  
 Ser Ser Thr Tyr Leu Arg Cys Asp Leu Thr Val Ser Met Gly Thr Glu  
 1810 1815 1820  
 Arg Leu Pro Gly Ser Trp Pro Tyr Val Tyr Leu Cys Glu Glu Ser Ser  
 1825 1830 1835 1840  
 Leu Cys Leu Phe Glu Pro Asp His Trp Thr Glu Ser Val Phe Pro Ser  
 1845 1850 1855  
 Phe Ser Gly Leu Phe Leu Ser Pro Lys Val Glu Arg Asp Glu Val Leu  
 1860 1865 1870  
 Ile Tyr Asn Ser Ser Cys Asn Ile Thr Met Glu Thr Glu Ala Glu Met  
 1875 1880 1885  
 Glu Cys Glu Met Pro Asn Gln Pro Ile Thr Ala Lys Ile Thr Glu Ile  
 1890 1895 1900  
 Gln Lys Ser Trp Gly Gln Asn Thr Gln Gly Asn Phe Ser Phe Gln Phe  
 1905 1910 1915 1920  
 Cys Arg Arg Trp Ser Arg Pro His Ser Trp Phe Pro Gln Arg Val Pro  
 1925 1930 1935  
 His Asp Gly Asp Ser Val Thr Val Glu Thr Gly His Leu Leu Leu Leu  
 1940 1945 1950  
 Asp Ala Asn Thr Ser Phe Leu Asn Ser Leu His Ile Lys Gly Gly Lys  
 1955 1960 1965  
 Leu Ile Phe Met Asp Pro Gly Pro Ile Glu Leu Arg Ala His Ser Ile  
 1970 1975 1980  
 Leu Ile Thr Asp Gly Gly Glu Leu His Ile Gly Ser Glu Glu Lys Pro  
 1985 1990 1995 2000  
 Phe Gln Gly Lys Ala Arg Ile Lys Ile Tyr Gly Ser Val His Ser Thr  
 2005 2010 2015  
 Pro Phe Phe Pro Tyr Gly Val Lys Phe Leu Ala Val Arg Asn Gly Thr  
 2020 2025 2030  
 Leu Ser Leu His Gly Ser Val Pro Glu Val Thr Val Thr Tyr Leu Gln  
 2035 2040 2045  
 Ala Ala Ala His Ala Gly Asp Lys Val Leu Thr Leu Gly Glu Ala Val  
 2050 2055 2060  
 Asp Trp Lys Pro Gly Asp Glu Ala Val Ile Thr Ser Gly Met Thr Val  
 2065 2070 2075 2080  
 Ala Gly Ala Glu Ala Thr Glu Val Val Val Glu Thr Val His Asn  
 2085 2090 2095  
 Ala Asp Leu His Leu Arg Asn Pro Leu Arg Tyr Ser Tyr Asp Phe Arg  
 2100 2105 2110  
 Glu Asn Trp Val Ala Gly Glu Asn Pro Ile Leu Lys Pro Thr Val Ala  
 2115 2120 2125  
 Leu Leu Ser Arg Asn Ile Ile Ile Gln Gly Asn Phe Thr Leu Glu Arg  
 2130 2135 2140  
 Val Lys Leu Leu Asn Ser Cys Gln Glu Ala Asn Thr Ala Lys Gly Asn  
 2145 2150 2155 2160  
 Leu Lys His Cys Leu Tyr Ser Lys Ser Glu Lys Met Leu Gly Ala Arg  
 2165 2170 2175  
 Asn Leu Gly Ala Arg Val Ile Ile Gln Ser Phe Pro Glu Glu Pro Ser  
 2180 2185 2190  
 Leu Val Lys Leu Lys Gly Val Gln Phe Arg Asp Leu Gly Gln Ala Phe  
 2195 2200 2205

His Lys His Leu Ser Ser Leu Thr Leu Val Gly Ala Met Arg Gly Ser  
 2210 2215 2220  
 Tyr Ile Gln Ser Cys Ser Val Trp Asn Ser Phe Ser Arg Gly Leu Ser  
 2225 2230 2235 2240  
 Met His Arg Thr Trp Gly Leu Lys Val Asp Ser Asn Val Phe Tyr Lys  
 2245 2250 2255  
 Ile Val Gly His Ala Leu Leu Leu Gly Ser Tyr Leu Asp Gly Arg Phe  
 2260 2265 2270  
 Ser Thr Ser Glu Thr Val Thr Gly Arg Lys Asn Gly Trp Trp Glu Gln  
 2275 2280 2285  
 Gly Ser Thr Ile Arg Asn Asn Val Ile Ile Ser Val Ser Ala Ala Glu  
 2290 2295 2300  
 Gly Leu Ser Gly Ser Glu Met Leu Ala Pro Ala Gly Ile Tyr Thr Phe  
 2305 2310 2315 2320  
 Ser Pro Thr Asn Val Met Glu Gly Asn Arg Val Cys Ala Ala Gly Tyr  
 2325 2330 2335  
 Gly Tyr Val Phe His Leu Val Thr Ser Gln Thr Leu Gln Ala Pro Leu  
 2340 2345 2350  
 Leu Ser Phe Asn Trp Asn Thr Ala His Ser Cys Thr Arg Tyr Gly Leu  
 2355 2360 2365  
 Leu Val Tyr Pro Lys Phe Gln Pro Pro Trp Asn Asn Asp Thr Gly Phe  
 2370 2375 2380  
 Thr Leu Phe Gln Asn Phe Met Val Trp Gly Ser Ala Gly Gly Ala Gln  
 2385 2390 2395 2400  
 Ile Phe Arg Ser Asn Asn Leu His Leu Lys Asn Phe Gln Val Tyr Ala  
 2405 2410 2415  
 Cys Arg Asp Phe Gly Ile Asp Ile Leu Glu Ser Asp Ala Asn Thr Leu  
 2420 2425 2430  
 Ile Thr Asp Ser Phe Leu Leu Gly His Phe Thr His Lys Gly Ser Leu  
 2435 2440 2445  
 Cys Met Ser Ala Gly Ile Lys Thr Pro Gln Arg Trp Glu Leu Thr Ile  
 2450 2455 2460  
 Ser Asn Thr Thr Phe Val Asn Phe Asp Gly Asn Cys Val Ala Ile Arg  
 2465 2470 2475 2480  
 Thr Cys Ser Gly Cys Phe Gln Gly Gln Gly Tyr Thr Val Lys Thr  
 2485 2490 2495  
 Arg Gln Leu Lys Phe Val Asn Ser Ser Asn Leu Val Ala Phe Xaa Phe  
 2500 2505 2510  
 Pro His Ala Ala Val Leu Glu Asp Leu Asp Gly Ser Leu Ser Gly Lys  
 2515 2520 2525  
 Asn Gly Ser His Val Leu Ala Ser Met Glu Thr Leu Ser Asp Thr Cys  
 2530 2535 2540  
 Leu Thr Asn Ala Ser Phe Ser Gln Ile Val Pro Gly Ser Val Cys Gly  
 2545 2550 2555 2560  
 Glu Ala Val Leu Phe His Arg Met Ser Ile Ala Leu Ala Asn Ser Leu  
 2565 2570 2575  
 Asp Val Pro Lys Asn Leu Thr Ile Thr Asp Ile Ser Asn Lys Thr Ile  
 2580 2585 2590  
 Thr Val Asn Tyr Val Glu Asp Thr Leu Ser Asn Tyr Tyr Gly Trp Met  
 2595 2600 2605  
 Ala Leu Leu Asp Gln Glu Thr Tyr Ser Leu Gln Phe Glu Ser Pro  
 2610 2615 2620

Trp	Met	Asn	Arg	Ser	Leu	Gln	Tyr	Ser	Ala	Thr	Phe	Asp	Ser	Phe	Ala
2625					2630				2635						2640
Pro	Gly	Asn	Tyr	Leu	Leu	Ile	Met	His	Arg	Asp	Leu	Pro	Pro	Tyr	Pro
				2645					2650					2655	
Asp	Ile	Leu	Leu	Arg	Cys	Gly	Ser	Gln	Val	Gly	His	Ser	Leu	Pro	Phe
			2660					2665					2670		
His	Pro	Leu	Pro	Ser	Gln	Asp	Arg	Ala	Cys	Asp	Trp	Phe	Phe	Asn	Arg
		2675					2680					2685			
Gln	Leu	Arg	Gln	Leu	Thr	Tyr	Leu	Val	Ser	Gly	Glu	Gly	Gln	Val	Lys
	2690					2695					2700				
Val	Phe	Leu	Gln	Leu	Lys	Pro	Gly	Val	Pro	Pro	Ser	Val	Ser	Ala	Ser
2705					2710					2715					2720
Thr	Ser	Val	Pro	Glu	Ser	Ala	Ser	Arg	Trp	Ser	Leu	Pro	Glu	Thr	Trp
				2725					2730					2735	
Gln	Asp	Val	Glu	Lys	Gly	Trp	Gly	Gly	Tyr	Asn	His	Thr	Ile	Pro	Gly
		2740						2745					2750		
Pro	Gly	Asp	Asp	Val	Leu	Ile	Leu	Pro	Asn	Lys	Thr	Val	Leu	Val	Asp
		2755					2760					2765			
Thr	Asp	Leu	Pro	Val	Leu	Arg	Cys	Leu	Tyr	Val	Met	Gly	Thr	Leu	Glu
	2770					2775					2780				
Phe	Pro	Val	Asp	Arg	Ser	Asn	Val	Leu	Ser	Val	Ala	Cys	Leu	Leu	Ile
2785					2790					2795					2800
Ala	Gly	Gly	Glu	Leu	Lys	Val	Gly	Thr	Leu	Glu	Asn	Pro	Leu	Glu	Lys
				2805					2810					2815	
Asp	Gln	Arg	Leu	Leu	Ile	Phe	Leu	Arg	Ala	Ser	Glu	Glu	Val	Val	Cys
			2820					2825					2830		
Asp	Tyr	Phe	Glu	Gly	Ile	His	Val	Asp	Pro	Gly	Thr	Ile	Gly	Val	Tyr
		2835					2840					2845			
Gly	Lys	Leu	Arg	Leu	His	Ser	Ala	Tyr	Pro	Lys	Lys	Ser	Trp	Val	His
	2850					2855					2860				
Leu	Gly	Ala	Asp	Ile	Ala	Pro	Gly	Asn	Glu	Arg	Ile	Ile	Val	His	Asn
2865					2870				2875						2880
Ala	Val	Asp	Trp	Gln	Pro	His	Asp	Thr	Ile	Val	Leu	Ser	Ser	Ser	Ser
				2885					2890					2895	
Tyr	Glu	Ala	His	Glu	Ala	Glu	Val	Leu	Thr	Val	Lys	Glu	Val	Lys	Gly
			2900					2905					2910		
His	His	Ile	Arg	Ile	Tyr	Glu	Arg	Leu	Lys	His	Arg	His	Ile	Gly	Ser
		2915					2920					2925			
Thr	His	Thr	Met	Glu	Asp	Gly	Gln	Gln	Val	His	Leu	Ala	Ala	Glu	Val
	2930					2935					2940				
Gly	Leu	Leu	Thr	Arg	Asn	Ile	Arg	Ile	Gln	Pro	Asp	Ser	Ser	Cys	Arg
2945					2950					2955					2960
Gly	Arg	Leu	Leu	Val	Gly	Ser	Phe	Arg	Lys	Ser	Ser	Gly	Glu	Asp	Phe
				2965											



Asn Gly His Gly Ile Asp Val Glu Gly Gln Asn Tyr Ser Leu Thr Asn  
 3045 3050 3055  
 Asn Leu Val Ile Leu Thr Met Gln Ser Ala Asn Ser Ser Pro Trp Val  
 3060 3065 3070  
 Ala Gly Ile Lys Val Asn Tyr Ala Glu Asp Ile Ile Leu His Gly Asn  
 3075 3080 3085  
 Val Val Ala Gly Ser Glu Arg Leu Gly Phe His Val Gly Gly His Gly  
 3090 3095 3100  
 Cys Ser Ser Glu Val Leu Trp Ser Asp Asn Val Val His Ser Ser Leu  
 3105 3110 3115 3120  
 His Gly Leu His Leu Tyr Lys Lys His Glu Ser Asn Asn Cys Thr Gly  
 3125 3130 3135  
 Val Ser Gly Phe Met Ala Phe Lys Asn Phe Asp Tyr Gly Ala Met Val  
 3140 3145 3150  
 Gln Thr Glu Asn Ser Val Asp Ile Gln Asn Ile Thr Leu Val Asp Asn  
 3155 3160 3165  
 Thr Val Gly Leu Leu Ala Ile Thr Tyr Val Ser Ser Ala Leu Leu Ser  
 3170 3175 3180  
 Ser Val Ser Thr Val Gln Ile Thr Leu Arg Asn Ser Val Ile Val Ala  
 3185 3190 3195 3200  
 Thr Ser Ser Ser Phe Asp Cys Ile His Asp Arg Lys Ala Pro Gln Ser  
 3205 3210 3215  
 Ala Asn Trp Thr Ser Thr Asp Arg Ala Pro Ser Asn Pro Arg Gly Gly  
 3220 3225 3230  
 Arg Ile Gly Ile Leu Trp Pro Val Ser Ala Ser Glu Pro Asn Ala Trp  
 3235 3240 3245  
 Pro Gln Glu Pro Trp His Lys Val Arg Ser Arg His Ser Val Pro Gly  
 3250 3255 3260  
 Ile Met Lys Leu Gln Asp Val Thr Phe Ser Ser Phe Val Lys Ser Cys  
 3265 3270 3275 3280  
 Tyr Ser Asn Asp Leu Asp Val Cys Ile Leu Pro Asn Glu Tyr Ser Thr  
 3285 3290 3295  
 Gly Val Met Tyr Pro Ile Thr Ala Glu Arg Thr Arg Met Leu Gly Ile  
 3300 3305 3310  
 Lys Asp Lys Asn Lys Phe Tyr Phe Pro Val Leu Gln Ser Ser Lys Asp  
 3315 3320 3325  
 Leu Val Gly Thr Ile Cys Pro Thr Leu Val Cys Glu Tyr Pro Arg Lys  
 3330 3335 3340  
 Tyr Leu Phe Thr Asp Leu Asp Gly Arg Thr Leu Gly Leu Pro Pro Pro  
 3345 3350 3355 3360  
 Val Ser Val Phe Pro Arg Thr Glu Glu Glu Trp Thr Gly Ser Phe Leu  
 3365 3370 3375  
 Asn Thr Gly Ile Phe Arg Glu Glu Gln Lys Cys Thr Phe Arg Ala Met  
 3380 3385 3390  
 Asn Gln Gly Phe Phe Cys Lys Gln Thr Glu His Ala Val Leu Ile Leu  
 3395 3400 3405  
 Asp Asn Val Asp Ala Thr Trp Thr Ile Pro Lys Ser His Pro Leu Val  
 3410 3415 3420  
 Ser Val Thr Asn Gly Phe Val Asp Thr Phe Ser Ile Val Lys Asp Ser  
 3425 3430 3435 3440  
 Asp Leu Cys Pro Pro Thr Ser Ser Leu Ser Thr Phe Tyr Ser Ile Leu  
 3445 3450 3455

Pro Thr Arg Gln Met Thr Lys Val Cys Phe Pro Glu Gln Thr Pro Pro  
 3460 3465 3470  
 Phe Leu Arg Phe Leu Leu Leu Gly Asn Gln Arg Ala Ser Lys Leu Ile  
 3475 3480 3485  
 Leu Ala Val Phe Tyr Asn Glu Ile Gln Ser Pro His Val Phe Leu Asp  
 3490 3495 3500  
 Lys Ser Phe Ile Pro Pro Thr Pro Leu Glu Ser Ala Phe Ser Leu Leu  
 3505 3510 3515 3520  
 Ala Glu Pro Ser Gly Ala Asn Tyr Phe Asp Ile Met Asn Asn Leu Leu  
 3525 3530 3535  
 Tyr Val Val Leu Gln Gly Glu Glu Pro Val Glu Ile His Ser Ser Val  
 3540 3545 3550  
 Ser Ile His Leu Ala Leu Thr Val Thr Phe Ser Val Leu Glu Lys Gly  
 3555 3560 3565  
 Trp Glu Arg Ala Met Leu Glu Ser Leu Ser Asp Phe Phe Gln Ile Asp  
 3570 3575 3580  
 Pro Asn Gln Ile Arg Leu Thr Leu Glu Met Pro Gly Asn Lys Glu Thr  
 3585 3590 3595 3600  
 Leu Glu Ala Ile Ala Asn Ser Glu Arg Lys Arg Lys Arg Asn Cys Pro  
 3605 3610 3615  
 Ser Val Thr Cys Gly Gly Pro Ser Ile Arg Tyr Gly Gln Arg Arg Pro  
 3620 3625 3630  
 Leu Met Ala Glu Met Thr Ser Leu Lys Ile Thr Pro Ala Thr Thr Leu  
 3635 3640 3645  
 Glu Thr Phe Ser Lys Val Ile Val Ile Glu Val Gly Asp Leu Pro Asn  
 3650 3655 3660  
 Ile Arg Asn Ser Glu Pro Ile Gln Ser Leu Pro Ser Asn Arg Leu Gln  
 3665 3670 3675 3680  
 Arg Leu Val Asn Gln Val Ile Thr Ala Gln Gln Thr Gly Ala Leu Glu  
 3685 3690 3695  
 Asn Val Leu Gly Met Thr Val Gly Ala Leu Leu Val Thr Gln Ser Lys  
 3700 3705 3710  
 Gly Val Thr Gly Tyr Arg Asn Ala Ser Ser Leu Ile Thr Gly Asn Leu  
 3715 3720 3725  
 Ile Tyr Thr Arg Pro Ser Glu Leu Ser Ile Leu Val Gln Pro Ser Asp  
 3730 3735 3740  
 Gly Glu Val Gly Ile Glu Leu Pro Val Gln Pro Arg Leu Val Phe Leu  
 3745 3750 3755 3760  
 Asp Glu Lys Asn Glu Arg Val Glu Ser Leu Gly Leu Pro Ser Glu Pro  
 3765 3770 3775  
 Trp Ile Ile Ser Val Ser Leu Glu Gly Ala Ser Glu Ser Val Leu Lys  
 3780 3785 3790  
 Gly Cys Thr Leu Ala Glu Thr Arg Asp Gly Tyr Val Thr Phe Ser Arg  
 3795 3800 3805  
 Leu Ala Val Leu Ile Ser Gly Ser Asn Trp His Leu Phe Phe Thr Val  
 3810 3815 3820  
 Ile Ser Pro Pro Gly Thr Asn Phe Thr Ala Arg Ser Arg Thr Phe Val  
 3825 3830 3835 3840  
 Val Leu Pro Val Ala Ser Lys Glu Arg Ser Thr Ile Ile Leu Ala Leu  
 3845 3850 3855  
 Ser Leu Cys Ser Val Ala Ser Trp Val Ala Leu Ser Cys Leu Val Cys  
 3860 3865 3870

Cys Trp Phe Lys Lys Ser Lys Thr Arg Lys Ile Lys Pro Glu Asp Ile  
 3875 3880 3885  
 Ser Glu Ser Gln Ala Lys Glu Gln Lys Lys Asn Thr His Asn Ser Ser  
 3890 3895 3900  
 Lys Pro Arg Gly Leu Gln Ala Lys Thr Ala Lys Glu Asn Thr Leu Met  
 3905 3910 3915 3920  
 Gly Glu Asp Met Arg Met Lys Val Met Gln Gly Met Gln Ser Gln Phe  
 3925 3930 3935  
 Pro Gln His Ser Met Asp Gly Val Ser Lys Arg Lys Val Ser Arg Leu  
 3940 3945 3950  
 Ala Val Thr Gly Glu Arg Thr Ala Thr Pro Ala Pro Lys Ile Pro Arg  
 3955 3960 3965  
 Ile Thr Cys Val Pro Gly Ser Leu Ala Gln Gln Leu Thr Leu Gln Glu  
 3970 3975 3980  
 Pro Gly Asn Trp Gln Glu Ala Gln Gln Gln Leu Leu Arg Tyr Gln Leu  
 3985 3990 3995 4000  
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 Arg Lys Gln Gly Gln Glu Pro Ser Gln Leu Asp Lys Gly Ser Asp Cys  
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<211> 4051

<212> PRT

<213> Rattus norvegicus

<400> 7

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 Ala Gly Gly Thr Trp Ile Thr Leu Val Phe Asp Gly Leu Glu Thr Ser  
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 Ile Leu Tyr Pro Asn Asn Gly Ser Gln Leu Ala Ile Asp Leu Val Asn  
 50 55 60  
 Val Ala Thr Pro Ala Leu Lys Ile Pro Cys Asp Ile Ser Pro Leu Phe  
 65 70 75 80  
 Val Asp Leu Pro Val Val Thr Cys Gln Thr Arg Ser Leu Pro Ser Glu  
 85 90 95  
 Ala Pro Glu Gly Leu Tyr Ser Leu Glu Ile Arg Ser Gly Asp Gln Leu  
 100 105 110  
 Leu Gly Ser Pro Cys Ala Arg Leu Leu Asp Asn Cys Thr Phe Lys Phe  
 115 120 125  
 Ser Lys Glu Gln Thr Pro Val Leu Tyr Gln Val Asn Pro Ser Ser Gly  
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 Val Pro Gly Glu Val Val His Val Tyr Gly Trp Val Ile Thr Asp Arg  
 145 150 155 160  
 Met Glu Ile Phe Asp Pro Asp Val Asp Tyr Met Glu Ser Pro Leu Ile

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Leu	Glu	Ala	Gly	Gly	Asp	Lys	Trp	Leu	Thr	Pro	Cys	Ser	Leu	Ile	Asn	
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Arg	Lys	Thr	Gly	Ile	Cys	Phe	Pro	Ile	Gln	Glu	Glu	His	Gly	Leu	Gly	
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Thr	Val	Gln	Cys	Arg	Val	Glu	Gly	Ser	Tyr	Ile	Gly	Ser	Gln	Asn	Val	
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Ser	Phe	Ser	Val	Phe	Asn	Lys	Gly	Arg	Ser	Met	Val	His	Lys	Glu	Ala	
225					230				235				240			
Trp	Leu	Ile	Ser	Ala	Lys	Gln	Glu	Leu	Phe	Leu	Tyr	Gln	Thr	Tyr	Pro	
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Asp	Ile	Thr	Ile	Thr	Gly	Asp	Phe	Asp	Pro	Ser	Ala	Arg	Val	Thr		
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Ile	Ala	Gly	Ile	Pro	Cys	Asp	Ile	Arg	His	Val	Ser	Pro	Lys	Lys	Ile	
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Glu	Cys	Thr	Thr	Arg	Ala	Pro	Arg	Asn	Gly	Ala	Arg	Leu	Thr	Ala	Pro	
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Gln	Ala	Ser	Asn	Arg	Gly	Leu	Leu	Phe	Glu	Val	Gly	Asn	Ala	Val	Lys	
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Tyr	Thr	Phe	Trp	Ile	Gln	Ala	Asp	Ser	Gln	Ala	Phe	Leu	Arg	Phe	Ser	
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Ser	Ser	Glu	Glu	Pro	Arg	Met	Lys	Val	Glu	Val	Ala	Ser	Ile	Ala	Val	
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Gly	Thr	Ala	Asp	Trp	Phe	Asp	Ser	Trp	Glu	Gln	Asn	Gly	Asn	Glu	Gly	
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Ser	Trp	Gln	Gln	Lys	Thr	Asn	Lys	Leu	Glu	Leu	Gln	Gly	Gly	Ala	Lys	
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Tyr	Tyr	Leu	Glu	Ala	Glu	Gln	His	Gly	Ile	Ala	Pro	Ser	Arg	Gly	Met	
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Arg	Ile	Ala	Val	Gln	Ile	His	Asn	Thr	Trp	Leu	Asn	Pro	Asp	Val	Val	
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Asn	Thr	Tyr	Leu	Leu	Glu	Lys	His	Gln	Ile	Arg	Ala	Gln	Ala	Gln	Arg	
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Leu	Pro	Glu	Ile	Gln	Val	Leu	Asn	Met	Ser	Gly	Lys	Gly	Asn	Phe	Phe	
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Ala	Gln	Gln	Ile	Gln	Thr	Ile	Ile	Glu	Glu	Leu	Leu	Val	Val	Lys	Cys	
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Asn	Leu	Val	Pro	Leu	Ser	Ala	Arg	Val	Leu	Leu	Trp	Leu	Gly	Phe	Glu	
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Gln	Gly	Leu	Asp	Gly	Ser	Arg	Ser	Asp	Gly	Val	Leu	Thr	Ser	Ser	Thr	
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Glu	Pro	Phe	Cys	Gly	Arg	Phe	Ser	Leu	Gly	Gln	Leu	Arg	His	Leu	Ile	

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Leu	Ser	Pro	Gly	Ala	Val	Ser	Lys	Gly	Tyr	Gln	Leu	Asp	Arg	Tyr	Pro		
595						600						605					
Tyr	Leu	Cys	Phe	Ala	Tyr	Arg	Gly	His	Met	Asn	Arg	Thr	Leu	Asp	Met		
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Thr	Val	Ser	Phe	Leu	Phe	Gly	Phe	Gln	Asn	Val	Met	Lys	Asn	Ile	Thr		
625			630						635			640					
Cys	Asp	Trp	Ser	Leu	Met	Glu	Pro	His	Pro	Glu	Ser	Trp	His	Phe	Thr		
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Cys	Asn	Asn	Leu	Trp	Asp	Thr	Cys	Val	Arg	His	Ser	Glu	Glu	Leu	Gln		
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Ser	Ser	Leu	Ala	Asn	Thr	Pro	Leu	Leu	Val	His	Arg	Ile	Asp	Ile	Phe		
675						680						685					
Pro	Val	Val	Pro	Glu	Ala	Asp	Leu	Leu	Tyr	Val	Asp	Glu	Ile	Ile	Leu		
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Pro	Gly	Gly	Asn	Val	Val	Glu	Ser	Val	Ser	Val	Val	Gly	Val	Pro	Pro		
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Val	Tyr	Ser	Ile	Ala	Ser	Trp	Met	Ala	Gly	Cys	Gly	Ser	Glu	Leu	Pro		
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Glu	Leu	Ile	Glu	Val	Thr	Ala	Lys	Arg	Leu	Gln	Arg	Thr	Ser	Pro	Pro		
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785			790						795			800					
Val	Pro	Val	His	Val	Ser	Ala	Arg	Gln	Leu	His	Lys	Leu	Leu	Gln	Asp		
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Asn	Ala	Asp	Glu	Ser	Thr	Ser	Gly	Tyr	Leu	Asn	Ala	Asp	Asp	Phe	Ile		
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Val	Thr	Lys	Asp	Gln	Asn	Ser	Cys	Tyr	Glu	His	Val	Trp	Thr	Leu	Ser		
835						840						845					
Trp	Thr	Thr	Gln	Ile	Gly	Asp	Leu	Pro	Asn	Phe	Ile	Arg	Val	Ser	Asp		
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Gln	Asn	Leu	Thr	Gly	Leu	Asn	Pro	Thr	Ile	Thr	Ala	Arg	Val	Val	Tyr		
865			870						875			880					
Asp	Gly	Gly	Val	Phe	Leu	Gly	Pro	Ile	Leu	Gly	Asp	Thr	Leu	Ala	Thr		
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Ala	Asn	Gln	Gln	Thr	Gln	Val	Thr	Val	Gln	Val	Asn	Asp	Ile	Pro	Ala		
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His	Cys	Ser	Gly	Ser	Cys	Ser	Phe	Gln	Tyr	Gln	Arg	Ala	Ser	Thr	Pro		
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Ser	Val	Asp	Cys	Val	Trp	Tyr	Ser	Leu	Gly	Ser	Asp	Val	Asn	Leu	Leu		
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Val	His	Phe	Thr	Gly	Asn	Gly	Phe	Pro	Arg	Asp	Ala	Gln	Phe	Leu	Gln		
945			950						955			960					
Val	Thr	Val	Ile	Asn	Thr	Ser	Cys	Glu	Val	Ile	Phe	Ser	Asn	Glu	Thr		
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Asn	Val	Ala	Cys	Glu	Leu	Ala	Leu	Leu	Pro	Val	Gly	Val	His	Arg	Val		
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Phe	Met	Leu	Val	Arg	Pro	Ser	Gly	Leu	Ala	Leu	Asn	Ala	Ser	Gly	Gly		

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Gly Leu Phe Leu Arg Val Glu Pro Arg Leu Asp Ala Val Glu Pro Ser		
1010	1015	1020
Thr Gly Ala Glu Ile Gly Gly Trp Trp Val Thr Leu Arg Gly Ser Ser		
1025	1030	1035
Leu Glu Gly Val Ser Leu Val Leu Phe Gly Thr Arg Ser Cys Val Ile		1040
1045	1050	1055
Asp Val Ile Arg Ser Asn Ser Gln Gln Ile Gln Cys Lys Val Pro Pro		
1060	1065	1070
Arg Gly Lys Asp Gly Tyr Thr Val Asn Val Thr Val Ile Asn Gly Asp		
1075	1080	1085
His Ser Thr Val Leu Ala Arg Ala Phe Thr Tyr Val Ser Ser Leu Asn		
1090	1095	1100
Pro Val Ile Val Ser Leu Ser Arg Asn Arg Ser Asn Ile Ala Gly Gly		
1105	1110	1115
Glu Ile Leu Phe Leu Gly Met Thr Leu Met Val Asn Tyr Thr Asp Leu		1120
1125	1130	1135
Asn Val Gln Ile Tyr Ile Gln Asn Thr Ser Ala Gln Val Leu Ala Lys		
1140	1145	1150
Thr Ala Trp Gly Leu Glu Val Leu Leu Pro Pro Leu Leu Pro Gly Ile		
1155	1160	1165
Tyr Val Ile Ser Ala Phe Ile Asn Gly Ile Ser Ile Arg Ser Gln Gly		
1170	1175	1180
Val Asp Leu His Ile Gln Tyr Leu Thr Glu Val Phe Ser Met Glu Pro		
1185	1190	1195
Cys Ser Gly Ser Leu Leu Gly Gly Thr Ile Leu Arg Leu Ser Gly Ala		1200
1205	1210	1215
Gly Leu Gly Arg Asp Pro Ala Leu Ile Arg Val Leu Val Asp Asn Arg		
1220	1225	1230
Pro Cys Asp Thr Val Lys Leu Thr Glu Val Asn Ile Trp Cys Glu Thr		
1235	1240	1245
Pro Arg Val Leu Leu Pro Pro Arg Ala Asp Val Leu Thr Val Pro Ala		
1250	1255	1260
Ser Val Glu Ile Trp Ala Gly Asn Thr Ser Phe Val Gly Lys Ala Phe		
1265	1270	1275
Ile Phe Thr Tyr Glu Ala Ala Ala Thr Pro Val Val Thr Ala Met Trp		1280
1285	1290	1295
Gly Glu Leu Met Asn Asn Ser Val Arg Phe Tyr Val Glu Gly Asn Asn		
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Leu Ser Asp Ser Val Ile Leu Leu Gly Ser Leu Lys Cys Asp Leu Gln		
1315	1320	1325
Val Gln Leu Phe Gly Asp Asn Met Asn Leu Ser Gly Cys Ser Leu Leu		
1330	1335	1340
Leu His Ser Leu Glu Ala Gly Val Tyr Ala Leu Gln Val Arg His Lys		
1345	1350	1355
Arg Met Gly Phe Ala Asn Met Ser Ala Val Pro Gln Lys Phe Glu Leu		1360
1365	1370	1375
Ser Pro Gln Ile Ile Asp Ile Phe Pro Thr His Gly Ser Ile Tyr Gly		
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Gly Thr Val Leu Thr Val Lys Gly Thr Ala Phe Arg Ser Arg Arg Arg		
1395	1400	1405
Ser Val His Val Asp Leu Ser Gly Pro Phe Thr Cys Val Ile Leu Ser		

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Leu Gly Asp His Thr Val Leu Cys Gln Thr Lys Phe Val Gly Asp Gln		
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Phe Ser Glu Thr Ser Leu Ala Leu Asn Ile Thr Val Leu Val Asn Gly		1440
	1445	1450
Leu Thr Ser Lys Cys Glu Gly Asn Cys Thr Leu Phe Leu Glu Glu Ala		1455
	1460	1465
Ala Thr Pro Ile Val Asp Ala Leu Thr Ile Ser Ile Ser Gly Ser Leu		1470
	1475	1480
Thr Thr Val Leu Met Arg Gly Gln Arg Leu Gly Ser Ala Gly Glu Pro		1485
	1490	1495
Thr Ala Phe Val Asp Gly Gln Leu Leu Cys His Thr Thr Phe Phe Asn		1500
1505	1510	1515
Thr Ser Tyr Val Ala Cys Gln Met Arg Asp Leu Ala Pro Gly Phe His		1520
	1525	1530
Tyr Leu Ser Ala Val Arg Thr Ser Ala Gly Tyr Ala Cys Leu Asn Ser		1535
	1540	1545
Val Ser Arg Asn Phe Phe Val Val Pro Gln Val Phe Asp Tyr Phe Pro		1550
	1555	1560
Lys Asp Phe Ser Ile His Gly Gly Ser Leu Leu Thr Ile Lys Gly Thr		1565
	1570	1575
Thr Leu Arg Gly Trp Lys Thr Thr Phe Val Tyr Val Gly Gln Gln Ala		1580
1585	1590	1595
Cys Leu Thr Val Asn Ile Ser Ser Glu Phe Ile Gln Cys Ile Val Pro		1600
	1605	1610
Ala Gly Asn Gly Ser Ala Ala Leu Glu Ile Asp Val Asp Gly Val Leu		1615
	1620	1625
Tyr His Met Gly Phe Val Asp Tyr Ser Ser Ile Phe Thr Pro Glu Leu		1630
	1635	1640
Leu Ser Val Ser Arg Ser His Asp Ile Leu Thr Phe Thr Val Ala Arg		1645
	1650	1655
Ile Ser Gly Ala Ala Asn Val Asp Ile Leu Ile Gly Thr Ser Pro Cys		1660
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Leu Asp Val Ala Gly Asn Arg Thr Ile Leu Arg Cys Met Val Pro Leu		1680
	1685	1690
Leu Pro Ala Gly Glu Tyr Ala Val Thr Gly Tyr Asp His Ser Arg Gly		1695
	1700	1705
Trp Val Ser Ser Thr Leu Ile Leu Val Leu Arg Ala Thr Val Thr Ser		1710
	1715	1720
Val Thr Glu Asn Tyr Gly Cys Leu Gly Gly Arg Leu Leu His Val Leu		1725
	1730	1735
Gly Ala Gly Phe Ser Pro Gly Asn Ile Ser Ala Ala Val Cys Gly Ala		1740
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Pro Cys Gln Val Leu Ala Asn Ala Thr Val Ser Ala Phe Ser Cys Leu		1760
	1765	1770
Val Leu Pro Leu Asp Ala Ser Leu Ala Phe Leu Cys Asn Leu Arg His		1775
	1780	1785
Ala Glu Asp Ser Cys Gly Val Ser Arg Pro Thr Tyr Leu Gln Cys Asp		1790
	1795	1800
Leu Thr Val Ser Val Gly Met Glu Arg Leu Pro Gly Ser Trp Pro Tyr		1805
	1810	1815
Val Tyr Leu Cys Glu Glu Ser Ser Gln Gly Leu Phe Val Pro Asp His		1820

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Gln Ile Phe Leu Ser	Phe Ser Gly Leu Phe Leu Ser	Pro Lys Val Glu				
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Arg Asp Glu Val Leu Ile Tyr Asn Ser Ser Cys Asn Ile Thr Met Glu						
	1860		1865		1870	
Thr Glu Ala Glu Met Glu Cys Gly Thr Pro Asn Gln Pro Ile Thr Ala						
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Lys Ile Thr Glu Ile Gln Lys Ser Arg Gly Gln Asn Thr Gln Gly Asn						
	1890		1895		1900	
Phe Ser Phe Gln Phe Cys Arg Arg Trp Ser Arg Pro His Ser Trp Phe						
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Pro His Arg Val Pro Gln Asp Gly Asp Asn Val Thr Val Glu Thr Ser						
	1925		1930		1935	
His Leu Leu Leu Leu Asp Ala Asn Thr Ser Leu Leu Asn Ser Leu His						
	1940		1945		1950	
Ile Lys Gly Gly Lys Leu Ile Phe Met Glu Pro Gly Pro Ile Asp Leu						
	1955		1960		1965	
Arg Ala His Ser Ile Leu Ile Thr Asp Gly Gly Glu Leu His Ile Gly						
	1970		1975		1980	
Ser Glu Asp Lys Pro Phe Gln Gly Lys Ala Arg Ile Lys Leu Tyr Gly						
1985	1990		1995		2000	
Ser Val His Ser Thr Ser Phe Phe Pro Tyr Gly Val Lys Phe Leu Ala						
	2005		2010		2015	
Val Arg Asn Gly Thr Leu Ser Leu His Gly Ser Val Pro Glu Val Thr						
	2020		2025		2030	
Phe Thr Tyr Leu Arg Ala Ala Ala Cys Ala Gly Asp Thr Val Leu Ala						
	2035		2040		2045	
Leu Glu Glu Ala Val Ala Trp His Pro Gly Asp Glu Ala Val Ile Ser						
	2050		2055		2060	
Ser Gly Ala Thr Val Glu Gly Ala Glu Ala Thr Glu Glu Val Val Val						
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Val Glu Thr Val His Asp Ala Asp Leu His Leu Arg Asn Pro Leu Arg						
	2085		2090		2095	
Tyr Ser Tyr Asn Phe Thr Glu Asn Trp Val Thr Gly Val Asn His Ile						
	2100		2105		2110	
Leu Lys Ala Thr Val Ala Leu Leu Ser Arg Asn Ile Ile Val Gln Gly						
	2115		2120		2125	
Asn Leu Thr Leu Glu Arg Val Lys Leu Leu Asp Ser Cys Gln Glu Ala						
	2130		2135		2140	
Gly Ala Ala Glu Gly Asn Leu Lys His Cys Leu Tyr Ser Lys Ser Glu						
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Lys Met Leu Gly Ala Arg Asn Leu Gly Ala Arg Val Ile Ile Gln Ser						
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Phe Pro Glu Glu Pro Ser Leu Val Lys Leu Lys Gly Val Gln Phe Arg						
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Asp Leu Gly Gln Ala Phe His Lys His Leu Ser Ser Leu Thr Leu Val						
	2195		2200		2205	
Gly Ala Met Arg Gly Ser Tyr Leu His Ser Cys Ser Met Trp Gly Ser						
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Phe Ser Arg Gly Leu Ser Ile His Arg Thr Trp Gly Leu Lys Val Glu						
2225	2230		2235		2240	
Asn Asn Ile Phe Tyr Lys Ile Val Gly His Ala Leu Leu Val Gly Ser						



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Tyr	Met	Asp	Arg	Ser	Phe	Ser	Thr	Ser	Glu	Ala	Val	Ile	Gly	Arg	Lys	
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Asn	Gly	Trp	Trp	Glu	Glu	Gly	Ser	Met	Ile	Arg	Asn	Asn	Val	Ile	Ile	
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Val	Cys	Ala	Ala	Gly	Tyr	Gly	Tyr	Val	Leu	His	Leu	Val	Thr	Ser	Gln	
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Thr	Val	Gln	Ala	Pro	Leu	Leu	Ser	Phe	Asn	Trp	Asn	Thr	Ala	His	Ser	
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Cys	Thr	Arg	Tyr	Gly	Leu	Leu	Val	Tyr	Pro	Glu	Phe	Gln	Pro	Pro	Trp	
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Thr	His	Glu	Gly	Ser	Leu	Cys	Met	Ser	Val	Gly	Ile	Lys	Thr	Pro	Lys	
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Gly	Phe	Pro	Val	Lys	Thr	Arg	Gln	Leu	Lys	Phe	Val	Asn	Ser	Ser	Asn	
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Leu	Val	Ala	Phe	Pro	Phe	Pro	His	Ala	Ala	Val	Leu	Gln	Asp	Leu	Asp	
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Thr	Leu	Ser	Asp	Thr	Cys	Leu	Thr	Asn	Ala	Ser	Phe	Ser	Gln	Ile	Val	
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Gly	Leu	Ala	Lys	Ser	Leu	Asp	Val	Pro	Lys	Asn	Leu	Thr	Met	Thr	Asp	
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Ile	Arg	Asn	Arg	Thr	Ile	Thr	Ile	Asn	Tyr	Val	Asp					

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Lys	Thr	Val	Leu	Val	Asp	Thr	Asp	Leu	Pro	Val	Leu	Arg	Cys	Leu	Tyr
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Val	Ala	Cys	Leu	Leu	Ile	Ala	Gly	Gly	Glu	Leu	Lys	Val	Gly	Thr	Leu
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Glu	Asn	Pro	Leu	Gly	Lys	Asp	Gln	Arg	Leu	Leu	Ile	Leu	Leu	Arg	Ala
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Ser	Glu	Glu	Ile	Phe	Cys	Asp	Arg	Phe	Asp	Gly	Ile	Arg	Val	Asp	Pro
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Lys	Lys	Ser	Trp	Val	His	Leu	Gly	Ala	Asp	Ile	Ala	Pro	Gly	Asn	Glu
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Arg	Ile	Ile	Val	His	Asn	Ala	Val	Asp	Trp	Gln	Pro	His	Asp	Lys	Ile
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Val	Leu	Ser	Ser	Ser	Ser	Tyr	Glu	Pro	His	Glu	Ala	Glu	Val	Leu	Thr
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Cys	Leu	Ala	Ala	Glu	Val	Gly	Leu	Leu	Thr	Arg	Asn	Ile	Arg	Ile	Gln
2930					2935					2940					
Ser	Asp	Ser	Ser	Cys	Arg	Gly	Arg	Leu	Leu	Val	Gly	Ser	Phe	Arg	Lys
2945	2950					2955					2960				
Ser	Ser	Gly	Glu	Glu	Phe	Ser	Gly	Ile	Leu	Gln	Leu	Leu	Asn	Val	Glu
2965					2970					2975					
Ile	Gln	Asn	Met	Gly	Ser	Pro	Leu	Tyr	Ala	Ser	Ile	Glu	Phe	Pro	Gly
2980					2985					2990					
Val	Ser	Ala	Gly	Ser	Trp	Val	Ile	Ser	Ser	Thr	Val	His	Gln	Ser	Cys
2995					3000					3005					
Thr	Val	Gly	Ile	His	Ala	Ser	Ser	Ser	Arg	Gly	Met	Val	Leu	Asn	Asp
3010					3015					3020					
Asn	Ile	Val	Phe	Gly	Thr	Lys	Gly	His	Gly	Ile	Asp	Val	Glu	Gly	Gln
3025	3030					3035					3040				
Asn	Tyr	Ser	Leu	Thr	Asn	Asn	Leu	Val	Ile	Leu	Ser	Met	Gln	Ser	Ala
3045					3050					3055					
Asn	Ser	Leu	Pro	Trp	Val	Ala	Gly	Ile	Lys	Val	Asn	Tyr	Ala	Glu	Asp
3060					3065					3070					
Ile	Ile	Leu	His	Gly	Asn	Val	Val	Ala	Gly	Ser	Glu	Arg	Leu	Gly	Phe

3075	3080	3085
His Val Arg Gly	His Gly Cys Ser Ser Glu Val	Leu Trp Ser Asp Asn
3090	3095	3100
Val Val His Ser Ser	Leu His Gly Ile His Leu Tyr Lys Lys Arg Glu	
3105	3110	3115
Pro Asn Asn Cys Thr Gly Val Ser Gly Phe Leu Ala Phe Lys Asn Phe		3120
3125	3130	3135
Asp Tyr Gly Ala Met Val Gln Thr Glu Asn Ser Val Asp Ile Gln Asn		
3140	3145	3150
Ile Thr Leu Val Asp Asn Thr Val Gly Leu Leu Ala Ile Ser Tyr Val		
3155	3160	3165
Ser Ser Ala Pro Leu Arg Ser Gly Ser Thr Leu Gln Ile Thr Leu Arg		
3170	3175	3180
Asn Ser Val Ile Val Ala Thr Ser Ser Ser Phe Asp Cys Ile Gln Asp		
3185	3190	3195
Arg Lys Ala Pro Gln Ser Ala Asn Trp Thr Ser Thr Asp Arg Ala Pro		3200
3205	3210	3215
Ser Asn Pro Arg Gly Gly Arg Ile Gly Ile Leu Trp Pro Val Ser Ala		
3220	3225	3230
Ser Glu Pro Asn Gly Trp Pro Gln Glu Pro Trp His Lys Val Arg Gly		
3235	3240	3245
Gly His Ser Val Pro Gly Ile Met Lys Leu Gln Asp Val Thr Phe Ser		
3250	3255	3260
Ser Phe Val Lys Ser Cys Tyr Ser Asp Asp Leu Asp Val Cys Ile Leu		
3265	3270	3275
Thr Asn Glu His Ser Thr Gly Ile Met Tyr Pro Ile Thr Ala Glu Arg		3280
3285	3290	3295
Thr Arg Met Leu Lys Ile Lys Asp Lys Asn Lys Phe Tyr Phe Leu Pro		
3300	3305	3310
Leu Gln Ser Gly Lys Asp Leu Val Gly Thr Ile Cys Pro Ala Ser Ser		
3315	3320	3325
Asp Cys Glu Ile Pro Arg Lys Tyr Leu Phe Thr Asp Leu Asp Gly Arg		
3330	3335	3340
Thr Leu Gly Leu Ser Pro Pro Val Ser Val Phe Pro Arg Ile Glu Glu		
3345	3350	3355
Gly Trp Thr Gly Ser Phe Phe Asn Thr Gly Ile Phe Arg Glu Glu Gln		3360
3365	3370	3375
Lys Cys Thr Phe Arg Ala Met Ile Gln Gly Phe Phe Cys Lys Gln Thr		
3380	3385	3390
Glu His Ala Val Leu Ile Leu Asp Asn Val Asp Val Thr Trp Thr Ile		
3395	3400	3405
Pro Lys Phe Tyr Pro Leu Val Ser Ile Thr Asn Gly Phe Val Asp Thr		
3410	3415	3420
Phe Ser Ile Val Lys Asp Ser Gly Leu Cys Phe Pro Thr Ser Ser Pro		
3425	3430	3435
Ser Thr Phe Tyr Ser Ile Leu Pro Thr Ser Gln Met Thr Lys Val Cys		3440
3445	3450	3455
Phe Pro Glu Arg Thr Pro Pro Phe Leu His Phe Phe Leu Leu Ala Asn		
3460	3465	3470
Arg Arg Ala Ser Lys Leu Ile Leu Ala Val Phe Tyr Asn Glu Ile Gln		
3475	3480	3485
Ser Pro Tyr Val Phe Leu Asp Lys Ser Phe Ile Pro Pro Thr Pro Val		

3490	3495	3500
Glu Ser Ala Phe Ser Leu Leu Asp Glu Pro Val Gly Ala Asn Tyr Phe		
3505	3510	3515 3520
Asp Ile Met Asn Asn Leu Leu Tyr Val Val Leu Gln Gly Glu Glu Ala		
	3525	3530 3535
Val Glu Ile Arg Ser Ser Val Ser Ile His Leu Ala Leu Thr Val Thr		
	3540	3545 3550
Phe Ser Val Leu Glu Lys Gly Trp Glu Lys Ala Val Leu Glu Ser Leu		
	3555	3560 3565
Arg Gly Phe Phe Gln Ile Asp Gln Asn Gln Ile Arg Leu Thr Leu Glu		
	3570	3575 3580
Met Pro Gly Asn Lys Glu Thr Leu Lys Ala Ile Ala Asn Ser Glu Gly		
3585	3590	3595 3600
Lys Arg Lys Arg Ser Cys Pro Thr Val Thr Cys Ala Val Pro Ser Ser		
	3605	3610 3615
Arg Tyr Val Gln Arg Arg Pro Leu Met Ala Glu Met Ala Ser Phe Thr		
	3620	3625 3630
Ile Thr Pro Ala Thr Thr Leu Glu Thr Phe Ser Lys Val Ile Val Phe		
	3635	3640 3645
Glu Ile Gly Asp Leu Pro Asn Ala Arg Asp Ser Glu Phe Ile Gln Ser		
	3650	3655 3660
Leu Pro Ser Asn Arg Leu Gln Lys Leu Ala His Gln Val Ile Thr Ala		
3665	3670	3675 3680
Gln Gln Thr Gly Val Leu Glu Asn Val Leu Gly Met Thr Val Gly Ala		
	3685	3690 3695
Leu Leu Val Ala Gln Ser Glu Gly Val Thr Gly Tyr Arg Asn Ala Ser		
	3700	3705 3710
Ser Leu Ile Thr Gly Asn Gln Ile Tyr Thr Arg Pro Ser Val Val Ser		
	3715	3720 3725
Ile Leu Val Gln Pro Ser Asp Gly Glu Val Gly Ile Glu Leu Pro Ile		
	3730	3735 3740
Gln Pro Arg Leu Ile Phe Leu Asp Glu Lys Asn Glu Arg Val Glu Ala		
3745	3750	3755 3760
Leu Gly Leu Pro Ser Glu Pro Trp Ile Ile Ser Val Ser Leu Glu Gly		
	3765	3770 3775
Thr Ser Glu Ser Met Leu Lys Gly Cys Thr Gln Ala Glu Thr Gln Asp		
	3780	3785 3790
Gly Tyr Val Thr Phe Ser Arg Leu Ala Val Leu Thr Ser Gly Ser Asn		
	3795	3800 3805
Trp His Leu Val Phe Thr Val Thr Phe Pro Pro Gly Thr Asn Leu Thr		
	3810	3815 3820
Ala Arg Ser Lys Thr Phe Ala Val Leu Pro Val Ala Gly Lys Glu Arg		
3825	3830	3835 3840
Ser Thr Ile Ile Leu Ala Leu Ser Leu Cys Ser Val Ala Ser Trp Val		
	3845	3850 3855
Ala Leu Ser Cys Leu Ile Cys Cys Trp Phe Lys Lys Ser Lys Thr Arg		
	3860	3865 3870
Lys Ile Lys Pro Glu Asp Ile Ser Glu Ser Gln Ala Lys Glu Gln Lys		
	3875	3880 3885
Lys Asn Thr His Ile Ser Ser Lys Pro Arg Glu Leu Gln Val Lys Thr		
	3890	3895 3900
Ala Lys Glu Asp Ala Met Met Gly Glu Asp Met Arg Met Lys Val Met		

3905                      3910                      3915                      3920  
 Glu Gly Arg Gln Asn Gln Phe Ser Gln His Ser Met Asp Gly Val Ser  
                                  3925                      3930                      3935  
 Lys Arg Lys Val Ser Arg Arg Ala Val Pro Glu Glu Arg Thr Ala Thr  
                                  3940                      3945                      3950  
 Pro Ala Leu Lys Ile Pro Arg Thr Thr Cys Val Pro Gly Ala Leu Pro  
                                  3955                      3960                      3965  
 Gln Gln Leu Thr Leu Pro Glu Pro Gly Asn Trp Gln Glu Ala Gln Gln  
                                  3970                      3975                      3980  
 Gln Leu Leu Arg Tyr Gln Leu Ala Gly Arg Asp Gln Leu Leu Leu Leu  
 3985                      3990                      3995                      4000  
 Cys Pro Asp Leu Arg Gln Glu Arg Gln Gln Gly Gln Glu Pro Ser Gln  
                                  4005                      4010                      4015  
 Leu Asn Lys Gly Ser Gly Arg Cys Met Gly Leu Ser Gln Glu Lys Ala  
                                  4020                      4025                      4030  
 Thr Cys Val Pro Thr Glu Thr Phe Cys Leu His Thr Ala Pro Pro Glu  
                                  4035                      4040                      4045  
 Thr Ile Gln  
                                  4050

<210> 8  
 <211> 77  
 <212> PRT  
 <213> Homo sapien

<400> 8  
 Ser Glu Ile Leu Ser Val Phe Pro Glu Thr Gly Ser Leu Gly Gly Arg  
 1                      5                      10                      15  
 Thr Asn Ile Thr Ile Thr Gly Asp Phe Phe Asp Asn Ser Ala Gln Val  
                                  20                      25                      30  
 Thr Ile Ala Gly Ile Pro Cys Asp Ile Arg His Val Ser Pro Arg Lys  
                                  35                      40                      45  
 Ile Glu Cys Thr Thr Arg Ala Pro Gly Lys Asp Val Arg Leu Thr Thr  
                                  50                      55                      60  
 Pro Gln Pro Gly Asn Arg Gly Leu Leu Phe Glu Val Gly  
 65                      70                      75

<210> 9  
 <211> 85  
 <212> PRT  
 <213> Homo sapien

<400> 9  
 Pro Cys Val His Ser Val Trp Tyr Ser Ile Asp Gly Asp Ile Asn Leu  
 1                      5                      10                      15  
 Met Ile Tyr Ile Thr Gly Thr Gly Phe Ser Gly Asp Ser Gln Phe Leu  
                                  20                      25                      30  
 Gln Val Thr Val Asn Lys Thr Ser Cys Lys Val Ile Phe Ser Asn Gln  
                                  35                      40                      45  
 Thr Asn Val Val Cys Gln Thr Asp Leu Leu Pro Val Gly Met His Arg  
                                  50                      55                      60  
 Ile Leu Met Leu Val Arg Pro Ser Gly Leu Ala Ile Ser Ala Thr Gly

65                      70                      75                      80  
Glu Asp Leu Phe Leu  
                        85

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<210> 10
<211> 85
<212> PRT
<213> Homo sapien
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[illegible]

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<210> 11
<211> 92
<212> PRT
<213> Homo sapien
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<400> 11															
Thr	Glu	Val	Phe	Ser	Ile	Glu	Pro	Cys	Cys	Gly	Ser	Leu	Leu	Gly	Gly
1				5					10					15	
Thr	Ile	Leu	Ser	Ile	Ser	Gly	Ile	Gly	Phe	Ser	Arg	Asp	Pro	Ala	Leu
			20					25					30		
Val	Trp	Val	Leu	Val	Gly	Asn	Arg	Ser	Cys	Asp	Ile	Val	Asn	Leu	Thr
		35				40						45			
Glu	Ala	Ser	Ile	Trp	Cys	Glu	Thr	Leu	Pro	Ala	Pro	Gln	Ile	Pro	Asp
	50				55						60				
Ala	Gly	Ala	Pro	Thr	Val	Pro	Ala	Ala	Val	Glu	Val	Trp	Ala	Gly	Asn
65					70					75					80
Arg	Phe	Phe	Ala	Arg	Gly	Pro	Ser	Pro	Ser	Leu	Val				
				85					90						

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<210> 12
<211> 91
<212> PRT
<213> Homo sapien
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<400> 12  
Pro Arg Ile Met Ala Ile Phe Pro Ser Gln Gly Ser Ala Cys Gly Gly  
1 5 10 15  
Thr Ile Leu Thr Val Arg Gly Leu Leu Asn Ser Arg Arg Ser  
20 25 30

Val Arg Val Asp Leu Ser Gly Pro Phe Thr Cys Val Ile Leu Ser Leu  
           35                          40                          45  
 Gly Asp His Thr Ile Leu Cys Gln Val Ser Leu Glu Gly Asp Pro Leu  
       50                          55                          60  
 Pro Gly Ala Ser Phe Ser Leu Asn Val Thr Val Leu Val Asn Gly Leu  
 65                          70                          75                          80  
 Thr Ser Glu Cys Gln Gly Asn Cys Thr Leu Phe  
                           85                          90

<210> 13  
 <211> 85  
 <212> PRT  
 <213> Homo sapien

<400> 13  
 Pro Val Met Asp Ala Leu Ser Thr Asn Thr Ser Gly Ser Leu Thr Thr  
 1                          5                          10                          15  
 Val Leu Ile Arg Gly Gln Arg Leu Ala Thr Thr Ala Asp Glu Pro Met  
           20                          25                          30  
 Val Phe Val Asp Asp Gln Leu Pro Cys Asn Val Thr Phe Phe Asn Ala  
           35                          40                          45  
 Ser His Val Val Cys Gln Thr Arg Asp Leu Ala Pro Gly Pro His Tyr  
       50                          55                          60  
 Leu Ser Val Phe Tyr Thr Arg Asn Gly Tyr Ala Cys Ser Gly Asn Val  
 65                          70                          75                          80  
 Ser Arg His Phe Tyr  
                           85

<210> 14  
 <211> 87  
 <212> PRT  
 <213> Homo sapien

<400> 14  
 Pro Gln Val Phe His Tyr Phe Pro Lys Asn Phe Ser Leu His Gly Gly  
 1                          5                          10                          15  
 Ser Leu Leu Thr Ile Glu Gly Thr Gly Leu Arg Gly Gln Asn Thr Thr  
           20                          25                          30  
 Ser Val Tyr Ile Asp Gln Gln Thr Cys Leu Thr Val Asn Ile Gly Ala  
           35                          40                          45  
 Glu Leu Ile Arg Cys Ile Val Pro Thr Gly Asn Gly Ser Val Ala Leu  
       50                          55                          60  
 Glu Ile Glu Val Asp Gly Leu Trp Tyr His Ile Gly Val Ile Gly Tyr  
 65                          70                          75                          80  
 Asn Lys Ala Phe Thr Pro Glu  
                           85

<210> 15  
 <211> 92  
 <212> PRT  
 <213> Homo sapien

<400> 15

His 1	Leu	Ser	Leu	His 5	Ile	Glu	Pro	Glu	Glu	Gly	Ser	Leu	Ala	Gly 15	Gly
Thr	Trp	Ile	Thr	Val	Ile	Phe	Asp	Gly	Leu	Glu	Leu	Gly	Val	Leu	Tyr
			20					25					30		
Pro	Asn	Asn	Gly	Ser	Gln	Leu	Glu	Ile	His	Leu	Val	Asn	Val	Asn	Met
		35					40					45			
Val	Val	Pro	Ala	Leu	Arg	Ser	Val	Pro	Cys	Asp	Val	Phe	Pro	Val	Phe
		50				55					60				
Leu	Asp	Leu	Pro	Val	Val	Thr	Cys	Arg	Thr	Arg	Ser	Val	Leu	Ser	Glu
65					70					75					80
Ala	His	Glu	Gly	Leu	Tyr	Phe	Leu	Glu	Ala	Tyr	Phe				
				85					90						

<210> 16

<211> 109

<212> PRT

<213> Homo sapien

<400> 16

Pro	Ile	Val	His	Gln	Val	Tyr	Pro	Pro	Ser	Gly	Val	Pro	Gly	Lys	Leu
1				5					10					15	
Ile	His	Val	Tyr	Gly	Trp	Ile	Ile	Thr	Gly	Arg	Leu	Glu	Thr	Phe	Asp
			20					25					30		
Phe	Asp	Ala	Glu	Tyr	Ile	Asp	Ser	Pro	Val	Ile	Leu	Glu	Ala	Gln	Gly
		35					40					45			
Asp	Lys	Trp	Val	Thr	Pro	Cys	Ser	Leu	Ile	Asn	Arg	Gln	Met	Gly	Ser
	50					55				60					
Cys	Tyr	Pro	Ile	Gln	Glu	Asp	His	Gly	Leu	Gly	Thr	Leu	Gln	Cys	His
65				70					75					80	
Val	Glu	Gly	Asp	Tyr	Ile	Gly	Ser	Gln	Asn	Val	Ser	Phe	Ser	Val	Phe
			85					90						95	
Asn	Lys	Gly	Lys	Ser	Met	Val	His	Lys	Lys	Ala	Trp	Leu			
			100					105							

<210> 17

<211> 84

<212> PRT

<213> Homo sapien

<400> 17

[illegible]



<210> 18  
 <211> 82  
 <212> PRT  
 <213> Homo sapien

<400> 18  
 Pro Val Val Thr Ala Met Gln Gly Glu Ile Thr Asn Ser Ser Leu Ser  
 1 5 10 15  
 Leu His Val Gly Gly Ser Asn Leu Ser Asn Ser Val Ile Leu Leu Gly  
 20 25 30  
 Asn Leu Asn Cys Asp Val Glu Thr Gln Ser Phe Gln Gly Asn Val Ser  
 35 40 45  
 Leu Ser Gly Cys Ser Ile Pro Leu His Ser Leu Glu Ala Gly Ile Tyr  
 50 55 60  
 Pro Leu Gln Val Arg Gln Lys Gln Met Gly Phe Ala Asn Met Ser Val  
 65 70 75 80  
 Val Leu

<210> 19  
 <211> 85  
 <212> PRT  
 <213> Homo sapien

<400> 19  
 Pro Glu Leu Ile Ser Ile Ser Gln Ser Asp Asp Ile Leu Thr Phe Ala  
 1 5 10 15  
 Val Ala Gln Ile Ser Gly Ala Ala Asn Ile Asp Ile Phe Ile Gly Met  
 20 25 30  
 Ser Pro Cys Val Gly Val Ser Gly Asn His Thr Val Leu Gln Cys Val  
 35 40 45  
 Val Pro Ser Leu Pro Ala Gly Glu Tyr His Val Arg Gly Tyr Asp Cys  
 50 55 60  
 Ile Arg Gly Trp Ala Ser Ser Ala Leu Val Phe Thr Ser Arg Val Ile  
 65 70 75 80  
 Ile Thr Ala Val Thr  
 85

<210> 20  
 <211> 81  
 <212> PRT  
 <213> Mus musculus

<400> 20  
 Pro Ala Val Ala Ser Val Ser Pro Thr Ser Gly Ser Ile Ala Gly Gly  
 1 5 10 15  
 Thr Thr Leu Met Ile Thr Gly Asn Gly Phe Ser Pro Gly Asn Thr Thr  
 20 25 30  
 Val Thr Val Gly Asp Gln Pro Cys Gln Ile Thr Phe Ile Ser Ser Ser  
 35 40 45

Glu Val Tyr Cys Ser Thr Pro Ala Gly Arg Ala Gly Thr Ala Asn Leu  
 50 55 60  
 Lys Ile Ser Val Asn Ala Ile Ile Tyr Pro Pro Leu Ser Phe Thr Tyr  
 65 70 75 80  
 Ala

<210> 21  
 <211> 83  
 <212> PRT  
 <213> Mus musculus

<400> 21  
 Pro Val Ile Thr Ser Ile Ser Pro Arg Tyr Gly Pro Gln Ala Gly Gly  
 1 5 10 15  
 Thr Leu Leu Thr Leu Thr Gly Lys Tyr Leu Asn Ser Gly Asn Ser Arg  
 20 25 30  
 His Ile Ser Ile Gly Gly Lys Thr Cys Thr Leu Lys Ser Val Ser Asp  
 35 40 45  
 Ser Ile Leu Glu Cys Tyr Thr Pro Ala Gln Thr Thr Ser Asp Glu Phe  
 50 55 60  
 Pro Val Lys Leu Lys Ile Asp Leu Ala Asn Arg Glu Thr Ser Ser Phe  
 65 70 75 80  
 Ser Tyr Arg

<210> 22  
 <211> 85  
 <212> PRT  
 <213> Mus musculus

<400> 22  
 Pro Thr Phe Tyr Arg Val Ser Pro Ser Arg Gly Pro Leu Ser Gly Gly  
 1 5 10 15  
 Thr Trp Ile Gly Ile Glu Gly Ser His Leu Asn Ala Gly Ser Asp Val  
 20 25 30  
 Ala Val Ser Ile Gly Gly Arg Pro Cys Ser Phe Ser Trp Arg Asn Ser  
 35 40 45  
 Arg Glu Ile Arg Cys Leu Thr Pro Pro Gly His Thr Pro Gly Ser Ala  
 50 55 60  
 Pro Ile Val Ile Asn Ile Asn Arg Ala Gln Leu Ser Asn Pro Glu Val  
 65 70 75 80  
 Lys Tyr Asn Tyr Thr  
 85

<210> 23  
 <211> 86  
 <212> PRT  
 <213> Homo sapien

<400> 23  
 Pro Val Leu Ile Ala Val Gln Pro Leu Phe Gly Pro Arg Ala Gly Gly

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      1           5           10           15
Thr Cys Leu Thr Leu Glu Gly Gln Ser Leu Ser Val Gly Thr Ser Arg
      20           25           30
Ala Val Leu Val Asn Gly Thr Glu Cys Leu Leu Ala Arg Val Ser Glu
      35           40           45
Gly Gln Leu Leu Cys Ala Thr Pro Pro Gly Ala Thr Val Ala Ser Val
      50           55           60
Pro Leu Ser Leu Gln Val Gly Gly Ala Gln Val Pro Gly Ser Trp Thr
      65           70           75           80
Phe Gln Tyr Arg Glu Asp
      85

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<210> 24  
 <211> 88  
 <212> PRT  
 <213> Artificial Sequence

<220>  
 <223> Consensus

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<400> 24
Pro Val Ile Thr Ser Ile Ser Pro Ser Ser Gly Pro Leu Ser Gly Gly
  1           5           10           15
Thr Glu Ile Thr Ile Thr Gly Ser Asn Leu Gly Ser Gly Ser Glu Asp
      20           25           30
Ile Lys Val Thr Phe Gly Gly Thr Glu Cys Asp Val Ile Ser Ser Ser
      35           40           45
Ser Ser Thr Gln Ile Val Cys Lys Thr Pro Pro Val Ala Gly Gly Pro
      50           55           60
Ser Pro Pro Val Val Thr Val Ser Leu Asp Gly Asp Gly Leu Ser Ser
      65           70           75           80
Ser Pro Val Thr Phe Thr Tyr Val
      85

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<210> 25  
 <211> 1776  
 <212> DNA  
 <213> Homo sapien

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<400> 25
atgactgcct ggctgatctc tctgatgagt attgaagtac tacttttggc agtacgtcac      60
ctgagtttac atattgaacc tgaagaaggt agccttgacg ggggaacgtg gatcacagtc      120
atTTTTgatg gtttgagatt ggggtgttctt taccccaaca atggctctca attggagata      180
cacctggtga acgtgaacat ggtggtgccc gcaactgcgga gtgttccttg tgacgtcttt      240
cctgttttct tggatttgcc tgtggtgaca tgccggacca gatctgtgct gtctgaagca      300
catgagggtc tgtacttcct ggaagcatac ttcgggggac agctggtgag cagtccaaat      360
ccaggaccac gagatagctg tactttcaag ttttccaagg cgcagacacc catcggtcac      420
caagtttatc caccaagtgg tgttcaggga aaactaatac atgtatatgg ctggattatc      480
actggaagat tggaaacttt tgattttgat gctgagtaca ttgatagccc agtgatcttg      540
gaagctcaag gagacaaatg ggttactcct tgctctctta taaataggca gatgggaagc      600
tgttatccta ttcaggagga ccatggtctt gggactctgc agtgccatgt ggaaggcgac      660
tacatcggtc cccagaatgt tagcttctca gtatttaaca aaggaaagtc aatggtccac      720

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aagaaggcat ggctgatcag tgctaaacag gatcttttcc tataccagac acactcagaa      780
atattatctg tgtttccaga aactggggagc cttgggggaa gaacaaacat cacaattaca      840
ggagactttt ttgacaattc tgcccagggtt accattgcag gcattccatg tgatattaga      900
cacgtgtctc ccaggaagat tgagtgcacc actcgggctc caggaaaaga tgtgaggctc      960
accacccctc agccaggcaa tcgagggctt ctttttgaag ttggagatgc tgttgaggga    1020
ctggaactga ctgaagccac cccagggtag aggtggcaga ttgtccctaa tgccagttct    1080
ccatttgggt tttggtcaca ggaaggacaa cctttcagag cacggctcag tgggttcttt    1140
gtggctccag agacaaataa ttacactttc tggattcagg cagatagcca agcttccttg    1200
catttcagtt ggtcagagga accaaggact aagggtgaaag tggcctccat cagcgtcggc    1260
actgctgact ggtttgactc ctgggagcag aatagggatg aagggacctg gcagcagaag    1320
actcccaagt tggagctgtt ggggtggagcc atgtactacc tggaagcaga gcatcatggg    1380
atagcccca gagggggatg gaggattggt gtccagattc acaacacctg gctgaatcct    1440
gatgtggtca ccacttacct acggggagaag caccagatcc gagtccgagc ccagaggctt    1500
ccagaagtac aggtgctgaa tgtatcaggc agaggaaact tcttccttac ttgggacaat    1560
gtctctagtc agccaatccc tgcaaatgcc acagcccatc tgattcaaac aaccattgag    1620
gagttacttg cagtaaaatg caaactggaa cccctttggt ctaacatcct tctccggtt    1680
ggatttgaac gaggcccaga agtttccaac tctgatgggg acctcaccag tgggacggag    1740
cccttctgtg gcaggttcag cctccgtcag cctcga      1776

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<210> 26

<211> 1871

<212> PRT

<213> Mus musculus

<400> 26

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Met Gly His Leu Trp Leu Ser Gly Thr Trp Phe Leu Phe Gly Leu Leu
 1           5           10          15
Trp Cys Ala Ala Asp Ser His Lys Gly Ser Ser Glu Thr Ile Pro Lys
 20          25          30
Val Thr Glu Val Ile Pro Lys Tyr Gly Ser Ile Asn Gly Ala Thr Arg
 35          40          45
Leu Thr Ile Lys Gly Glu Gly Phe Ser Gln Ala Ser Gln Phe Asn Tyr
 50          55          60
Gly Ala Asp Asn Thr Glu Leu Gly Asn His Val Gln Leu Val Ser Ser
 65          70          75          80
Phe Gln Ser Ile Thr Cys Asp Val Glu Lys Asp Ser Ser His Ser Thr
 85          90          95
Gln Ile Thr Cys Tyr Thr Arg Ala Met Pro Glu Asp Thr Tyr Ser Val
100         105         110
Arg Val Ser Val Asp Gly Val Pro Val Ala Glu Asn Asn Thr Cys Lys
115         120         125
Gly Val Ala Ser Ser Trp Ala Cys Ser Phe Ser Thr Lys Ser Phe Arg
130         135         140
Thr Pro Thr Ile Arg Ser Ile Thr Pro Leu Ser Gly Thr Pro Gly Thr
145         150         155         160
Leu Ile Thr Ile Lys Gly Arg Leu Phe Thr Asp Val Tyr Gly Ser Asn
165         170         175
Thr Ala Leu Ser Ser Asn Gly Arg Asn Val Arg Ile Leu Arg Ile Tyr
180         185         190
Ile Gly Gly Met Pro Cys Glu Leu Leu Ile Pro His Ser Asp Asp Leu
195         200         205
Tyr Gly Leu Lys Leu Asp His Ala Asn Gly Asp Thr Gly Ser Val Thr

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210		215		220
Cys Lys Thr Thr Gly	Thr Tyr Ile Gly His His Asn Val Ser Phe Ile			
225	230	235		240
Leu Asp Ser Asp Tyr Gly Arg Ser Phe Pro Glu Lys Met Thr Tyr Phe				
	245	250		255
Val Ser Ser Leu Asn Lys Ile Ser Met Phe Gln Thr Tyr Pro Glu Val				
	260	265		270
Val Met Val Ser Pro Ser Lys Gly Ser Thr Glu Gly Gly Thr Leu Leu				
	275	280		285
Thr Ile His Gly His Phe Phe Asp Gln Thr Asp Leu Pro Val Arg Val				
	290	295		300
Leu Val Gly Gly Gln Ala Cys Ala Ile Leu Asn Val Thr Glu Asn Thr				
305	310	315		320
Ile Tyr Cys Lys Thr Pro Pro Lys Pro His Ile Leu Lys Ala Thr Tyr				
	325	330		335
Pro Gly Gly Arg Gly Leu Lys Val Glu Val Trp Asn Asn Ser Arg Pro				
	340	345		350
Ala His Leu Glu Asp Ile Leu Glu Tyr Asn Glu His Thr Pro Gly Tyr				
	355	360		365
Met Gly Ala Thr Trp Thr Asp Ser Ala Ser Tyr Val Trp Pro Ile Glu				
	370	375		380
Gln Gly Thr Phe Val Ala Arg Ile Ser Gly Phe Leu Val Pro Pro Asp				
385	390	395		400
Ser Asp Val Tyr Arg Phe Tyr Ile Arg Gly Asp Asp Arg Tyr Ala Ile				
	405	410		415
Tyr Phe Ser Gln Thr Gly Arg Thr Glu Asp Lys Val Arg Ile Ala Tyr				
	420	425		430
Tyr Ser Gly Asn Ala Asn Thr Tyr Phe Ser Asn Ser Thr Gln Arg Ser				
	435	440		445
Asp Glu Ile His Leu Gln Lys Gly Lys Glu Tyr Tyr Ile Glu Ile Leu				
	450	455		460
Leu Gln Glu Tyr Thr Leu Ser Ala Phe Val Asp Val Gly Leu Tyr Gln				
465	470	475		480
Tyr Lys Asn Val Phe Thr Glu Gln Gln Thr Gly Asp Ala Leu Asn Glu				
	485	490		495
Glu Gln Val Ile Lys Ser Gln Ser Thr Val Val Pro Glu Val Gln Ile				
	500	505		510
Ile Thr Leu Glu Asn Trp Glu Thr Ala Asp Val Thr Asn Glu Val Gln				
	515	520		525
Gln Val Thr Val Thr Ser Pro Cys Val Gly Ala Asn Ser Cys Ser Leu				
	530	535		540
Ser Gln Tyr Arg Phe Ile Tyr Asn Met Glu Lys Thr Val Trp Leu Pro				
545	550	555		560
Ala Asp Ala Ser Asp Phe Thr Leu Lys Ser Ala Leu Asn Asp Leu Trp				
	565	570		575
Ser Ile Lys Pro Asp Ser Val Gln Val Thr Ser Lys Arg Asp Leu Gln				
	580	585		590
Ser Tyr Ile Tyr Thr Ile Thr Phe Val Ser Thr Arg Gly Asp Phe Asp				
	595	600		605
Leu Leu Gly Tyr Glu Val Phe Glu Gly Ser Asn Val Thr Leu Ser Ile				
	610	615		620
Thr Glu Gln Thr Lys Gly Lys Pro Asn Leu Glu Thr Phe Thr Leu Asn				

625	630										635				640	
Trp	Asp	Gly	Ile	Ala	Ser	Lys	Pro	Leu	Thr	Pro	Glu	Ser	Ser	Glu	Ala	
				645					650					655		
Glu	Phe	Gln	Val	Ala	Val	Glu	Glu	Met	Val	Ser	Ala	Lys	Cys	Pro	Pro	
			660					665					670			
Glu	Ile	Ala	His	Leu	Glu	Glu	Gly	Phe	Leu	Val	Lys	Tyr	Phe	Arg	Asp	
		675					680					685				
Tyr	Glu	Thr	Asp	Phe	Glu	Leu	Glu	His	Ile	Asn	Arg	Gly	Gln	Lys	Thr	
	690					695				700						
Ala	Glu	Thr	Asp	Ala	Tyr	Cys	Gly	Arg	Tyr	Ser	Leu	Lys	Asn	Pro	Ala	
705					710					715					720	
Val	Leu	Phe	Asp	Ser	Thr	Asp	Val	Lys	Pro	Asn	Lys	Ser	Pro	Tyr	Gly	
				725					730					735		
Asp	Ile	Leu	Leu	Phe	Pro	Tyr	Asn	Gln	Leu	Cys	Leu	Ala	Tyr	Lys	Gly	
		740						745					750			
Ser	Leu	Ala	Asn	Phe	Ile	Asp	Leu	Lys	Phe	Lys	Tyr	Gln	Asp	Ser	Gly	
		755					760					765				
Lys	Ile	Ile	Arg	Ser	Ala	Asp	Val	Gln	Phe	Glu	Tyr	Asn	Phe	Ala	Ser	
	770					775					780					
Gly	Asn	Lys	Trp	Thr	Tyr	Thr	Cys	Ile	Asp	Leu	Leu	Asp	Phe	Leu	Gln	
785					790					795					800	
Thr	Lys	Tyr	Ala	Gly	Thr	Ser	Phe	Ser	Leu	Gln	Arg	Ile	Thr	Leu	Gln	
				805					810					815		
Lys	Ser	Ser	Glu	Phe	Gln	Ser	Ile	Tyr	Val	Asp	Ala	Val	Tyr	Ile	Gly	
			820					825					830			
Gln	Thr	Pro	Thr	Val	Ser	Val	Leu	Asp	Asp	Met	Pro	Lys	Arg	Arg	Pro	
		835					840					845				
Pro	Ala	Leu	Ala	Asn	Lys	Gly	Ile	Phe	Leu	Lys	His	Phe	Gln	Val	Asn	
	850					855					860					
Arg	Thr	Lys	Leu	Asn	Gly	Ser	Ala	Met	Thr	Ile	Gln	Tyr	Ser	Val	Thr	
865					870					875					880	
Ile	Thr	Ser	Tyr	Asn	Cys	Ser	His	Asn	Ile	Pro	Met	Met	Ala	Val	Ser	
				885					890					895		
Phe	Gly	Gln	Ile	Ile	Thr	Asn	Glu	Thr	Lys	Asn	Glu	Leu	Val	Tyr	Arg	
			900					905					910			
Gly	Asn	Asn	Trp	Pro	Gly	Glu	Ser	Lys	Ile	Arg	Ile	Gln	Lys	Ile	Gln	
		915					920					925				
Glu	Ala	Ser	Pro	Pro	Ile	Ser	Gly	Ser	Phe	Asp	Val	Gln	Ala	Tyr	Gly	
	930					935					940					
His	Thr	Leu	Lys	Gly	Ile	Pro	Ala	Ala	Val	Pro	Ala	Ala	Asp	Leu	Gln	
945					950					955					960	
Phe	Ala	Leu	Gln	Ser	Leu	Glu	Glu	Ile	Glu	Gln	Val	Ser	Val	Asn	Arg	
				965					970					975		
Glu	Gly	Thr	Cys	Ala	Gly	Tyr	Ser	Trp	Ser	Ile	Arg	Trp	Thr	Ser	Pro	
		980						985								

	1045		1050		1055
Cys Gly Phe Thr Trp Asp Ala Met Ile Thr Pro Leu Ile Leu Thr Thr					
	1060		1065		1070
Thr Pro Ser Glu Gly Ser Tyr Ala Glu Ser Thr Ile Leu Thr Ile Ala					
	1075		1080		1085
Gly Ser Gly Phe Ser Pro Thr Ser Ala Val Ser Val Ser Val Gly Ser					
	1090		1095		1100
Thr Arg Cys Ser Leu Leu Ser Val Glu Glu Asn Glu Ile Lys Cys Gln					
1105		1110		1115	1120
Ile Val Asn Gly Ser Ala Gly His Val Pro Val Ala Val Ser Ile Ala					
	1125		1130		1135
Asp Val Gly Leu Ala Gln Asn Leu Glu Gly Glu Gly Ser His Phe Ile					
	1140		1145		1150
Tyr Arg Ser Gln Ile Ser His Val Trp Pro Asp Ser Gly Ser Leu Ala					
	1155		1160		1165
Gly Gly Thr Leu Leu Thr Ile Ser Gly Phe Gly Phe Ser Glu Asn Ser					
	1170		1175		1180
Thr Val Leu Val Gly Asn Glu Thr Cys Asn Val Ile Glu Gly Asp Leu					
1185		1190		1195	1200
Asn Arg Ile Thr Cys Arg Thr Ser Lys Arg Ile Glu Gly Thr Val Asp					
	1205		1210		1215
Ile Ser Val Ile Thr Asn Gly Ile Gln Val Thr Ala Lys Asp Ser Phe					
	1220		1225		1230
Ser Tyr Ser Cys Leu Gln Thr Pro Val Val Thr Asp Phe Ser Pro Lys					
	1235		1240		1245
Glu Arg Thr Val Leu Gly Lys Val Asn Leu Thr Ile Lys Gly Tyr Asn					
	1250		1255		1260
Phe Gly Asn Glu Leu Ala Gln Asn Thr Val Tyr Val Gly Arg Lys His					
1265		1270		1275	1280
Cys Gln Val Leu His Ser Asn Phe Thr Asp Ile Thr Cys Leu Leu Pro					
	1285		1290		1295
Thr Leu Pro Pro Gly Lys His Asp Ile Tyr Val Lys Val Arg Asn Trp					
	1300		1305		1310
Gly Leu Ala Ser Thr Arg Asn Lys Leu Asn Ala Ser Ile Leu Tyr Ile					
	1315		1320		1325
Leu Glu Val Ile His Met Phe Pro Gln Arg Gly Ser Leu Tyr Gly Gly					
	1330		1335		1340
Thr Glu Ile Thr Ile Met Gly Phe Gly Phe Ser Thr Ile Pro Thr Glu					
1345		1350		1355	1360
Asn Ser Val Leu Leu Gly Ser Phe Pro Cys Asp Ile Thr Ser Ser Ser					
	1365		1370		1375
Glu Asn Val Ile Lys Cys Thr Leu His Ser Thr Gly Thr Val Phe Arg					
	1380		1385		1390
Ile Thr Asn Asn Gly Ser His Leu Val His Gly Leu Gly Tyr Ala Trp					
	1395		1400		1405
Ser Pro Ser Val Leu Asn Val Thr Val Gly Asp Thr Val Val Trp Ser					
	1410		1415		1420
Trp Gln Ala His Pro Phe Leu Arg Gly Ile Gly Tyr Arg Ile Phe Ser					
1425		1430		1435	1440
Val Ser Ser Pro Gly Ser Val Thr Tyr Asp Asp Lys Gly Phe Thr Asn					
	1445		1450		1455
Gly Arg Gln Lys Ser Ala Ser Gly Ser Phe Ser Tyr Gln Phe Thr Ser					

1460	1465	1470
Pro Gly Ile Tyr Tyr Tyr Ser Ser Gly Tyr Val Asp Glu Ala His Ser		
1475	1480	1485
Ile Ser Leu Gln Gly Val Ile Asn Val Phe Pro Ala Glu Ala Arg His		
1490	1495	1500
Ile Pro Leu Tyr Leu Phe Val Gly Asn Ile Glu Ala Thr Tyr Val Pro		
1505	1510	1515
Ala Gly Pro Ala His Leu Gln Leu Ala Ser Thr Ala Ala Gly Cys Leu		
1525	1530	1535
Ala Thr Glu Pro Leu Cys Gly Leu Asn Asp Thr Arg Val Lys His Ser		
1540	1545	1550
Asn Lys Leu Phe Phe Glu Leu Ser Asn Cys Ile Ser Pro Ser Ile Ile		
1555	1560	1565
Asn Ile Thr Pro Ser Thr Gly Thr Ala Asn Glu Leu Ile Thr Ile Ile		
1570	1575	1580
Gly His Gly Phe Ser Ser Leu Pro Cys Ala Asn Lys Val Thr Ile Gly		
1585	1590	1595
Ser Tyr Pro Cys Val Val Glu Glu Ser Ser Glu Asn Ser Ile Ile Cys		
1605	1610	1615
His Ile Asp Pro Gln Asn Ser Met Asn Val Gly Ile Arg Glu Ile Val		
1620	1625	1630
Thr Leu Ile Val Tyr Asn Leu Gly Thr Ala Ile Asn Thr Leu Thr Lys		
1635	1640	1645
Ala Phe Asp Arg Arg Phe Val Leu Leu Pro Asn Ile Asp Met Val Met		
1650	1655	1660
Pro Lys Ala Gly Ser Thr Thr Gly Met Thr Arg Val Thr Ile Gln Gly		
1665	1670	1675
Ser Gly Phe Met Ser Ser Pro Glu Gly Val Glu Val Phe Met Gly Asp		
1685	1690	1695
Phe Pro Cys Lys Val Leu Ser Val Thr Tyr Thr Ala Leu Asn Val Glu		
1700	1705	1710
Thr Ser Pro Ala Pro Gln Gln Leu Val Leu Val Asp Ile Leu Ile His		
1715	1720	1725
Gly Val Pro Ala Gln Cys Gln Ser Asn Cys Ser Phe Ser Tyr Leu Glu		
1730	1735	1740
Asn Ile Ala Pro Tyr Val Thr Gly Ile Phe Pro Asn Ser Ile Gln Gly		
1745	1750	1755
Tyr Gly Asn Val Leu Ile Lys Gly Glu Arg Phe Gly Thr Val Leu Glu		
1765	1770	1775
Glu Ile Ser Ile Phe Ile Gly Ser Gln Gln Phe Arg Val Ile Asp Val		
1780	1785	1790
Asn Glu Asn Asn Ile Thr Val Leu Met Thr Pro Leu Glu Ala Gly Leu		
1795	1800	1805
His Ser Leu Ser Val Val Val Gly Ser Lys Gly Leu Ala Leu Gly Asn		
1810	1815	1820
Leu Thr Ile Ser Ser Pro Ala Val Ala Ser Val Ser Pro Thr Ser Gly		
1825	1830	1835
Ser Ile Ala Gly Gly Thr Thr Leu Met Ile Thr Gly Asn Gly Phe Ser		
1845	1850	1855
Pro Gly Asn Thr Thr Val Thr Val Gly Asp Gln Pro Cys Gln Ile		
1860	1865	1870



<210> 27  
 <211> 446  
 <212> DNA  
 <213> Homo sapien

<400> 27  
 tgtgactgga gtctcacgag gaccagcccc gagagctggc agttcgattg cactgacctc 60  
 tgggagactt gtgtgcggtt cttcggggat ctccagcccc ctccggcaaa ctccccagtg 120  
 ctgggttcac agatcaacct tctccctctg gccaggaga cgggcctgtt ctatgtggat 180  
 gaaattatta ttgcagacac aaacgtaaca gtttctcaag ctgattctgg aacggctcgc 240  
 ccagggggca atctggtgga atcagtctct gtggtgggat cccctccggt ctacagtgtc 300  
 acctcctggc tggcgggggtg tggcacggag ctcccgtca tcaactgcacg ctctgtgccc 360  
 actgaaggaa cagaagaggg atctggactg gtcctggtga cgacacagag acgacagcgg 420  
 acaagtccac ctctaggagg acactt 446

<210> 28  
 <211> 188  
 <212> DNA  
 <213> Homo sapien

<400> 28  
 tcttgacggt tacagtgaac aaaacgaggt gcaaagttat tttctcaaac cagaccaatg 60  
 tagtctgtca gacagatttg ctacctgttg gaatgcatcg gatcttgatg ttggtgagac 120  
 cctctggtct tgccatcagt gccactggag aagacctctt cctaaatgtg aaacctagac 180  
 tggatatg 188

<210> 29  
 <211> 116  
 <212> PRT  
 <213> Homo sapien

<220>  
 <221> VARIANT  
 <222> (1)...(433)  
 <223> Xaa = Any Amino Acid

<400> 29  
 Arg Leu Arg Asn Trp Asp Pro Gly Gln Asp Ser Ala Lys Gln Val Val  
 1 5 10 15  
 Ile Lys Glu Gly Asp Met Leu Arg Leu Thr Ser Asp Ala Thr Val His  
 20 25 30  
 Ser Ile Val Ile Gln Asp Gly Gly Leu Leu Val Phe Gly Asp Asn Lys  
 35 40 45  
 Asp Gly Ser Arg Asn Ile Thr Leu Arg Thr His Tyr Ile Leu Ile Gln  
 50 55 60  
 Asp Gly Gly Ala Leu His Ile Gly Ala Glu Lys Cys Arg Tyr Lys Ser  
 65 70 75 80  
 Lys Ala Thr Ile Thr Leu Tyr Gly Lys Ser Asp Glu Gly Glu Ser Met  
 85 90 95  
 Pro Thr Phe Gly Lys Lys Phe Ile Gly Val Glu Ala Gly Gly Thr Leu  
 100 105 110  
 Glu Leu His Gly

115

<210> 30  
 <211> 114  
 <212> PRT  
 <213> Homo sapien

<220>  
 <221> VARIANT  
 <222> (1)...(433)  
 <223> Xaa = Any Amino Acid

<400> 30  
 Glu Leu Gln Pro Trp Asn Pro Gly His Asp Gln Asp His His Val His  
 1 5 10 15  
 Ile Gly Gln Gly Lys Thr Leu Leu Leu Thr Ser Ser Ala Thr Val Tyr  
 20 25 30  
 Ser Ile His Ile Ser Glu Gly Gly Lys Leu Val Ile Lys Asp His Asp  
 35 40 45  
 Glu Pro Ile Val Leu Arg Thr Arg His Ile Leu Ile Asp Asn Gly Gly  
 50 55 60  
 Glu Leu His Ala Gly Ser Ala Leu Cys Pro Phe Gln Gly Asn Phe Thr  
 65 70 75 80  
 Ile Ile Leu Tyr Gly Arg Ala Asp Glu Gly Ile Gln Pro Asp Pro Tyr  
 85 90 95  
 Tyr Gly Leu Lys Tyr Ile Gly Val Gly Lys Gly Gly Ala Leu Glu Leu  
 100 105 110  
 His Gly

<210> 31  
 <211> 489  
 <212> DNA  
 <213> Homo sapien

<400> 31  
 cacagtcggc ttggggcctg gaggtggcac tgccccact gccagctggt ctccacagaa 60  
 tttccgtctc tatcaatggg gtcagcattc actcacaagg ggttgatctc cacatccagt 120  
 acctcacaga agttttcagc atcgagcctt gctgtgggtc cctgctggga gggaccatcc 180  
 tcagcatctc aggaataggc ttcagcaggg acccagcttt ggtttgggta cttgtgggca 240  
 atcggtcctg tgacattgtg aacttaacgg aggcgagcat ctggtgtgaa accctgccag 300  
 cccccagat acccgatgcg ggcgctccca ctgttccagc tgccgtggag gtctgggctg 360  
 gcaacagggt cttcgcccgt ggtccttcac caagcttggt ggggaaaggc ttcaccttca 420  
 tgtatgaagc ggcagcaaca ccagtagtca ctgccatgca aggagaaatc acaaatagca 480  
 gcctgagcc 489

<210> 32  
 <211> 366  
 <212> PRT  
 <213> Homo sapien

<400> 32

Ser	Ile	Val	Ala	Leu	Asn	Lys	Ser	Tyr	Glu	Val	Tyr	Phe	Thr	Gly	Thr	1	5	10	15
Ser	Pro	Gln	Asn	Leu	Arg	Leu	Met	Leu	Leu	Asn	Val	Asp	His	Asn	Lys	20	25	30	
Ala	Val	Leu	Val	Gly	Ile	Phe	Phe	Ser	Thr	Leu	Gln	Arg	Leu	Asp	Val	35	40	45	
Tyr	Val	Asn	Asn	Leu	Leu	Val	Cys	Pro	Lys	Thr	Thr	Ile	Trp	Asn	Ala	50	55	60	
Gln	Gln	Lys	His	Cys	Glu	Leu	Asn	Asn	His	Leu	Tyr	Lys	Asp	Gln	Phe	65	70	75	80
Leu	Pro	Asn	Leu	Asp	Ser	Thr	Val	Leu	Gly	Glu	Asn	Tyr	Phe	Asp	Gly	85	90	95	
Thr	Tyr	Gln	Met	Leu	Tyr	Leu	Leu	Val	Lys	Gly	Thr	Ile	Pro	Val	Glu	100	105	110	
Ile	His	Thr	Ala	Thr	Val	Ile	Phe	Val	Ser	Phe	Gln	Leu	Ser	Val	Ala	115	120	125	
Thr	Glu	Asp	Asp	Phe	Tyr	Thr	Ser	His	Asn	Leu	Val	Lys	Asn	Leu	Ala	130	135	140	
Leu	Phe	Leu	Lys	Ile	Pro	Ser	Asp	Lys	Ile	Arg	Ile	Ser	Lys	Ile	Arg	145	150	155	160
Gly	Lys	Ser	Leu	Arg	Arg	Lys	Arg	Ser	Met	Gly	Phe	Ile	Ile	Glu	Ile	165	170	175	
Glu	Ile	Gly	Asp	Pro	Pro	Ile	Gln	Phe	Ile	Ser	Asn	Gly	Thr	Thr	Gly	180	185	190	
Gln	Met	Gln	Leu	Ser	Glu	Leu	Gln	Glu	Ile	Ala	Gly	Ser	Leu	Gly	Gln	195	200	205	
Ala	Val	Ile	Leu	Gly	Asn	Ile	Ser	Ser	Ile	Leu	Gly	Phe	Asn	Ile	Ser	210	215	220	
Ser	Met	Ser	Ile	Thr	Asn	Pro	Leu	Pro	Ser	Pro	Ser	Asp	Ser	Gly	Trp	225	230	235	240
Ile	Asn	Val	Thr	Ala	Gln	Pro	Val	Glu	Arg	Ser	Ala	Phe	Pro	Val	His	245	250	255	
His	Val	Ala	Phe	Val	Ser	Ser	Leu	Leu	Val	Ile	Thr	Gln	Pro	Val	Ala	260	265	270	
Ala	Gln	Pro	Gly	Gln	Pro	Phe	Pro	Gln	Gln	Pro	Ser	Val	Lys	Ala	Thr	275	280	285	
Asp	Ser	Asp	Gly	Asn	Cys	Val	Ser	Val	Gly	Ile	Thr	Ala	Leu	Thr	Leu	290	295	300	
Arg	Ala	Ile	Leu	Lys	Asp	Ser	Asn	Asn	Asn	Gln	Val	Asn	Gly	Leu	Ser	305	310	315	320
Gly	Asn	Thr	Thr	Ile	Pro	Phe	Ser	Ser	Cys	Trp	Ala	Asn	Tyr	Thr	Asp	325	330	335	
Leu	Thr	Pro	Leu	Arg	Thr	Gly	Lys	Asn	Tyr	Lys	Ile	Glu	Phe	Ile	Leu	340	345	350	
Asp	Asn	Val	Val	Gly	Val	Glu	Ser	Arg	Thr	Phe	Ser	Leu	Leu			355	360	365	

&lt;210&gt; 33

&lt;211&gt; 30

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

<220>

<223> Primer

<400> 33

gccttccttggt ggaccattga ctttcctttg

30

<210> 34

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 34

gccagcattt ccgagtgggtg

20

<210> 35

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 35

tgggtaagca aggtcaaact tcc

23

<210> 36

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 36

accctcaata caagaaggaa ccac

24

<210> 37

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 37

ggtgctcccg tctcccagcg

20

<210> 38

<211> 18

<212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer

<400> 38  
 ggtctttgaa agaaggaa

18

<210> 39  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer

<400> 39  
 caatcaattc aaaggtctca g

21

<210> 40  
 <211> 22  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer

<400> 40  
 gcaggactac agaaatactc ag

22

<210> 41  
 <211> 20  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer

<400> 41  
 tcccacttac cacagagaag

20

<210> 42  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer

<400> 42  
 aggaggaaca tggatcacag t

21

<210> 43  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer

<400> 43  
 agccattggtt ggggtaaaga a 21

<210> 44  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer

<400> 44  
 ttataatcat gcaaaatggg c 21

<210> 45  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer

<400> 45  
 acatcatttt catgaggcaa t 21

<210> 46  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer

<400> 46  
 ggctgtgagg aatggaactc t 21

<210> 47  
 <211> 23  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer

<400> 47  
 atggtcattg tcctcatatt tgc 23  
  
 <210> 48  
 <211> 25  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Primer  
  
 <400> 48  
 aggtttcaga acagcaaaat aatcg 25  
  
 <210> 49  
 <211> 27  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Primer  
  
 <400> 49  
 ttctcaaggt aacctattgt gttctta 27  
  
 <210> 50  
 <211> 25  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Primer  
  
 <400> 50  
 tggtttgaat ctgaccttca aaacc 25  
  
 <210> 51  
 <211> 23  
 <212> DNA  
 <213> Artificial Sequence  
  
 <220>  
 <223> Primer  
  
 <400> 51  
 aaatgtgcac ttggtaaaac ccc 23  
  
 <210> 52  
 <211> 25  
 <212> DNA  
 <213> Artificial Sequence

<220>

<223> Primer

<400> 52

tttcacactg tcctgtgtca atgac

25

<210> 53

<211> 23

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 53

aaaatccctc atcctgtctg gtc

23

<210> 54

<211> 24

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 54

ttgggaattc atggtttttg attc

24

<210> 55

<211> 25

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 55

acataccttc ctccagcctt agaac

25

<210> 56

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Primer

<400> 56

gaaaggcttg tgcctcctgt g

21

<210> 57



<211> 25  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer

<400> 57  
 tggcaaacag attcacaatt attcc

25

<210> 58  
 <211> 21  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer

<400> 58  
 cattgagttt gagctaagtc c

21

<210> 59  
 <211> 22  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer

<400> 59  
 catgcagcat gtatgtaact ag

22

<210> 60  
 <211> 18  
 <212> DNA  
 <213> Artificial Sequence

<220>  
 <223> Primer

<400> 60  
 gtttattggg agtattgc

18

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&lt;213&gt; Homo sapien

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&lt;213&gt; Homo sapien

&lt;400&gt; 213

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&lt;211&gt; 2303

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&lt;213&gt; Homo sapien

&lt;400&gt; 214

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 Val Val Ala Ser Thr Asp Tyr Ser Met Tyr Gln Ala Glu Glu Phe Thr  
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 Leu Leu Pro Cys Ser Glu Cys Ser His Phe Gln Val Lys Val Lys Glu  
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 Thr Pro Gln Phe Leu His Met Gly Glu Ile Ile Asp Gly Val Asp Met  
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 Arg Ala Glu Val Gly Ile Leu Thr Arg Asn Ile Val Ile Gln Gly Glu  
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[illegible]

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Lys	Pro	Met	Tyr	Leu	His	Ile	Gly	Glu	Glu	Ile	Asp	Gly	Val	Asp	Met
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Arg	Gly	Gly	Tyr	Asp	Pro	Pro	Thr	Tyr	Ile	Arg	Asp	Leu	Ser	Ile	His
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His	Thr	Phe	Ser	Arg	Cys	Val	Thr	Val	His	Gly	Ser	Asn	Gly	Leu	Leu
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Ile	Lys	Asp	Val	Val	Gly	Tyr	Asn	Ser	Leu	Gly	His	Cys	Phe	Phe	Thr
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